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Result
No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Minimum DB
Maximum DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Total number of hits satisfying chosen parameters:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OM protein - protein search, using sw model
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                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
  448
448
375.5
351
347
334.5
                                                                                                                                                                                                                                                                                                                                           Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         seq
                                                                            A Geneseq_16Dec04:*
1: geneseqp1990s:*
2: geneseqp1990s:*
3: geneseqp2000s:*
4: geneseqp2001s:*
5: geneseqp2003s:*
6: geneseqp2003as:*
7: geneseqp2003as:*
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length: 2000000000
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Gapop 10.0 , Gapext 0.
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524
1 SMGGVPGAIPGGVPGGVFYP.....
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  Maximum Match 100%
Listing first 45 s
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Copyright (c) 1993 - 2005 Compugen Ltd.
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              AAR56653
AAY09135
AAY69135
AAY69071
AAY69071
AAY69069
AAY69069
AAY69069
AAW46315
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AAW1966637
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Aay01302 Human tro
Aay69069 Amino aci
Ade40112 Human NOV
Aab08630 Amino aci
Aaw46315 Human ela
Aab08631 Fusion pr
Aao17360 Human ela
Ad019747 Human ela
Ad019747 Human ela
Ad019747 Human ela
Ad0196420 Human ela
Abu08725 Human ela
Abu08725 Human ela
Abu08725 Human ela
Abu08725 Human ela
Abu08726 Human pro
Ad066216 Novel human bum
Ad064761 Human pro
Aab8422 Human mad
Ad068871 Novel pro
Adm03792 Human mad
                                                                                                                                                                                                                                                                                                                                           Description
                                                                                                                                                                                                                                                              Aay69071 Amino
Ade40134 Human
                                                                                                                                                                                                                                                                                       Aay01301 Amino
Aay69135 Amino
                                                                                                                                                                                                                                                                                                                 Aar56653 Synthetic
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aci
NOV
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	45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26
	173	173	173	173	173	173	173	173	173	174	174	174	174	174	174	174	177.5	201.5	267.5	326.5
	33.0	33.0						33.0				33.2						38.5		62.3
	1002	936	936	378	378	378	378	378	312	937	936	936	936	936	832	832	479	91	660	864
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ALIGNMENTS	AAW09218	ABP53472	AAR80251	ABW01635	ABG31419	AAY51889	AAW53545	AAW09219	ABP53480	AAW53547	ABW01637	ABG31421	AAY51891	AAW09221	ABP53473	AAR80252	AAW49739	AAU91258	AAY01303	ADE56670
	Aaw09218	Abp53472	Aar80251	Abw01635	Abg31419	Aay51889		Aaw09219	Abp53480		Abw01637	Abg31421		Aaw09221		Aar80252	Aaw49739	Aau91258	Aay01303	_
	SELPUK PO	Protein p	Polymer S	Plasmid p	SELPOK pr	Plasmid p	Amino aci	SELPOK po	Protein p	Amino aci	Plasmid p	SELPOK-CS	Plasmid p	SELPOK-CS	Protein p	Polymer S	Protein p	Nucleic a	Human tro	Rat Prote

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RESULT 1
AAR56653
Synthetic
                                                                                                                                                                               Synthetic human tropoelastin (SHEL).
                                                                                                                                                                                                                      AAR56653 standard; protein;
                                                                                 Weiss AS, Martin SL;
                                                                                                        22-DEC-1992;
28-JUN-1993;
                                                                                                                         16-DEC-1993;
                                                                                                                                    07-JUL-1994.
                                                                                                                                              WO9414958-A1.
                                                                                                                                                                   Tropoelastin; pharmaceutical; surgical dressing
                                                                                                                                                                                          25-MAR-2003
22-MAR-1995
                                                                                                                                                                                                           AAR56653;
                                                                                             (UNSY ) UNIV SYDNEY.
                                                                                                                                                                                          (revised)
(first entry)
                                                                                                       92AU-00006520.
93AU-00009661.
                                                                                                                         93WO-AU000655
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Human synthetic tropoelastin is susceptible to hydrolytic breakdown of the crosslinks. Such material may be useful in e.g. surgical applications, where the gradual loss of material over time is intended. (Updated on 25-MAR-2003 to correct PN field.)

Disclosure; Page 30; 77pp; English.

variants.

Synthetic polynucleotide(s) - encode recombinant tropoelastins

and

WPI; 1994-263633/32. N-PSDB; AAQ70941.

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                                                                                                                       Matches 100;
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Best Local :
                                                                                                                                                                                                          used to produce the proteins recombinantly. The tropoelastin derivatives or hybrid proteins containing the derivatives are useful in medical, pharmaceutical, veterinary and cosmetic applications, e.g. as antiwrinkle or hand lotions, also as surgical implants, foods and industrial products. The hybrid protein have controllable GAG-binding properties, depending on presence or absence of a specific fragment, designated peptide 26A, from hTE. The present sequence represents the amino acid sequence of the synthetic human tropoelastin SHEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY01301 standard; protein; 733
                                                                                                                                                                                Sequence 733 AA;
                                                                                                                                                                                                                                                                                                                                   The invention relates to a derivative or variant of human tropoelastin (hTE) having elastin-like and/or macromolecule (specifically glycosaminoglycan (GAG))-binding properties. Cells containing vectors comprising the nucleic acids encoding the variants or derivatives are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-132162/11.
N-PSDB; AAX27704.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tropoelastin; hTB; elastin; glycosaminoglycan; GAG-binding; medical; pharmaceutical; veterinary; cosmetic application; anti-wrinkle; food; hand lotion; surgical implant; industrial product; human; SHEL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Weiss AS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Amino acid sequence of synthetic human tropoelastin SHEL.
                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Fig 1; 82pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                          macromolecular binding properties, useful e.g. as surgical implants
                                                                                                                                                                                                                                                                                                                                                                                                                                                             New derivatives of human tropoelastin - with elastin-like or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (UNSY ) UNIV SYDNEY.
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                                                                                                                                    Similarity
                  VTFPGALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLGVS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  VTFPGALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLGVS
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                                                             SMGGVPGAIPGGVPGGVFYPGAGLGALGGGALGPGGKPLKPVPGGLAGAGLGAGLGAFPA
                                                                              SMGGVPGAIPGGVPGGVFYPGAGLGALGGGALGPGGKPLKPVPGGLAGAGLGAGLGAFPA
VTFPGALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLGVS
                                                                                                                       Conservative
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                                                                                                                                   100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                             English.
                                                                                                                       0
                                                                                                                   Score 524; Db 2;
Pred. No. 7.1e-36;
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Best Local S
Matches 98
                                                                                                                                                                                                                                              derivatives, and other polypeptides containing tropoclastin derivative-derived protease-susceptibility sites, are useful in human or veterinary medicine, cosmetics (e.g. antiwrinkle or hand lotions), as bulking agents and for inducing chemotaxis. They are also useful for proliferation or growth inhibition, particularly of smooth muscle cells, epithelial or endothelial cells, fibroblasts, osteocytes, chondrocytes and platelets. Peptidomimetics that mimic the protease cleavage site in tropoclastin derivatives are competitive inhibitors of the protease, and are used for protecting against lung damage caused by elastin, for inhibiting or controlling localized growth of cancers or metastases, or to limit protease activity that causes blood clotting
                                                                                                                                                                                                                                                                                                                                                                                                                                                 representative of troppelastin derivatives of the invention. In the troppelastin derivatives of the invention a subsequence has been mutated so that susceptibility to proteolysis is reduced or eliminated, or a subsequence has been inserted so that susceptibility to proteolysis is increased. The derivatives have with reduced susceptibility, and can be used where the wild-type protein would be degraded too easily, e.g. in contact with serum or wound exudate. The troppelastin derivatives provide competitive inhibition of protease activity. The troppelastin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hand lotion; bulking agent; chemotaxis; proliferation; growth inhibition; peptidomimetic; lung damage; elastin; cancer; metastasis; blood clotting.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tropoelastin;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY69135;
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                                                                                                                                                                                                                  Sequence 515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 131-133; 136pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-182399/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Amino acid sequence of a human tropoelastin derivative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence represents a human tropoelastin derivative,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New derivatives of tropoelastin useful, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UNSY ) UNIV SYDNEY
                                  63
61 FPGALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLGVS 98
                                                                      ...
                                                                                                                                             98;
                                                                                                                                                              Similarity
                                  FPGALVPGGVADAAAYKAAKAGAGLGGVPGVGGLGVS 100
                                                                    GGVPGA I PGGVPGGVFYPGAGLGALGGGALGPGGKPLKPVPGGLAGAGLGAGLGAFPAVT
                                                                                        GGVPGAIPGGVPGGVFYPGAGLGALGGGALGPGGKPLKPVPGGLAGAGLGAGLGAGLGAFPAVT 62
                                                                                                                                          98.3%; So larity 100.0%; 1 Conservative 0;
                                                                                                                                                                                                                   A
A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98AU-00004723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99WO-AU000580
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                                                                                                                                                              Score 515; DB 3;
Pred. No. 2.9e-35;
                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               for inducing chemotaxis
                                                                                                                                               ٥,
                                                                                                                                                                              Length 515;
                                                                                                                                             Indels
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                                                                                                                                               Gaps
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RESULT 4 AAY69071

ADE40134;

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RESULT 5
ADE40134
ID ADE4
XX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC subsequence has been inserted so that susceptibility to proteclysis is concreased. The derivatives have with reduced susceptibility, and can be conteact with serum or wound exudate. The tropoelastin derivatives provide conteact with serum or wound exudate. The tropoelastin derivatives provide competitive inhibition of protease activity. The tropoelastin derivatives containing tropoelastin derivatives and other polypeptides containing tropoelastin derivative. CC derivatives, and other polypeptides containing tropoelastin derivative and for inducing chemotaxis. They are also useful in human or veterinary cc and for inducing chemotaxis. They are also useful for proliferation or cc growth inhibition, particularly of smooth muscle cells, epithelial or cc endothelial cells, fibroblasts, osteocytes, chondrocytes and platelets. Cc peptidomimetics that mimic the protease cleavage site in tropoelastin cc derivatives are competitive inhibitors of the protease, and are used for controlling localized growth of cancers or metastases, or to limit
                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence represents a human tropoelastin derivative, which is representative of tropoelastin derivatives of the invention. In the tropoelastin derivatives of the invention a subsequence has been mutated so that susceptibility to proteolysis is reduced or eliminated, or a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tropoelastin; derivat hand lotion; bulking peptidomimetic; lung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 571 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New derivatives of tropoelastin useful, e.g. for inducing chemotaxis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAY69071 standard; protein; 571 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             activity that causes blood
                                                                                                                                                                                                                     FPGALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLGVS 100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 115-117; 136pp; English.
                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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bulking agent; chemotaxis; proliferation; growth inhibition;
c; lung damage; elastin; cancer; metastasis; blood clotting.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98AU-00004723.
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                                                                                                                                                                                                                                                                                                                                                                                                            98.3%;
100.0%;
                                                                                                                                                                                                                                                                                                                                                                             Score 515; Db s; Pred. No. 3.2e-35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tropoelastin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             clotting
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 571;
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                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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ADE40134 standard;

protein;

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The invention relates to a novel NOVX polypeptide. The invention demonstrates cardiant, antiarteriosclerotic, cytostatic, anorectic, antidiabetic, immunosuppressive

antiparkinsonian,

immunosuppressive, anti insonian, antiasthmatic

The polypeptide of the tic, hypotensive,

anti-HIV,

Claim 1;

SEQ

ID NO

40;

560pp;

English

New NOVX polypeptides and nucleic acids, useful for preventing or treating NOVX-associated disorders, e.g. cancer, cardiomyopathy, atherosclerosis or diabetes, and in chromosome mapping, tissue ty

tissue typing

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WPI; 2003-663472/62. N-PSDB; ADE40133.

Kekuda R, M Shimkets RA,

Guo X,

da R, Miller CE, Patturajan M, Pena CEA, Rieger DK; kete RA, Zerhusen BD, Li L, Ji W, Padigaru M, Casman SJ; EZ, Boldog FL, Gorman L, Leite MW, Vernet CAM, Anderson X, Zhong M, Gerlach VL, Hjalt T, Rastelli L, Spytek KA; ger SR, Ellerman K, Malyankar UM, Macdougall JR, Stone DJ; brook JP, Lepley DM, Burgess CE, Majumder K, Wolenc AR;

DW.

Alsobrook JP,

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02-AUG-2001;

03-AUG-2001;

07-AUG-2001;

08-AUG-2001;

09-AUG-2001;

13-AUG-2001;

16-AUG-2001;

17-AUG-2001;

20-AUG-2001;

20-AUG-2001;

20-AUG-2001;

21-AUG-2001;

22-AUG-2001;

23-AUG-2001;

23-AUG-2001;

23-AUG-2001;

24-AUG-2001;

25-AUG-2001;

27-AUG-2001;

28-AUG-2001;

29-AUG-2001;

29-AUG-2001;

19-APR-2002;

19-APR-2002;

19-APR-2002;
                                                                                                                                                                                                                                                          19-APR-2002;
23-APR-2002;
07-JUN-2002;
01-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NOVX; cardiant; antiarteriosclerotic; hypotensive; cytostatic; anorectic; antidiabetic; immunosuppressive; anti-HIV; neuroprotective; nootropic; antiparkinsonian; antiasthmatic; gynaecological; cardiomyopathy; atherosclerosis; hypertension; cancer; obesity; diabetes; AIDS; multiple sclerosis; graft-versus-host disease; Alzheimer's; Parkinson's; multiple sclerosis; graft-versus-host disease; Alzheimer's; Parkinson's;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-JAN-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      fertility disorder; vaccine; gene therapy;
                                                                                                                                                                                                                                                          2001US-0309501P
2001US-0310291P
2001US-0310951P
2001US-0311999P
2001US-0312892P
2001US-0313415P
2001US-0313415P
2001US-0313415P
2001US-0313415P
2001US-031443P
2001US-031403P
2001US-031403P
2001US-031403P
2001US-031403P
2001US-0315403P
2001US-0315403P
2001US-0322716P
2001US-0322716P
2001US-03493P
2002US-037494P
2002US-037494P
2002US-037494P
2002US-037493P
2002US-0373825P
2002US-0373825P
2002US-0373899P
2002US-03786971P
2002US-0386971P
2002US-0386971P
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RESULT 6
AAYO1302
ID AAYO
AC AAYO
AC AAYO
DT 07-J
XX Trop
KW Phaz
KW Hand
OS Synt
XX WO95
PN W095
PN W095
PN W095
AC AYO
DT 17-(
XX WAN
PT NEW
PT MACI
XX OC ON
CC ON
CC ON
CC ON
CC OPTOI
CC PPTO
CC PTOI
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Best Local :
The invention relates to a derivative or variant of human tropoelastin (hTE) having elastin-like and/or macromolecule (specifically glycosaminoglycan (GAG))-binding properties. Cells containing vectors comprising the nucleic acids encoding the variants or derivatives are used to produce the proteins recombinantly. The tropoelastin derivatives or hybrid proteins containing the derivatives are useful in medical, pharmaceutical, veterinary and cosmetic applications, e.g. as antivrinkle or hand lotions, also as surgical implants, foods and industrial products. The hybrid protein have controllable GAG-binding properties, depending on presence or absence of a specific fragment, designated peptide 26A, from hTE. The present sequence represents the synthetic human tropoelastin variant SHELdelta26A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gynaecological activities and may be useful in diagnosing, treating or preventing NOVX-associated disorders including cardiomyopathy, atherosclerosis, hypertension, cancer, catchest, and a parkinson's sclerosis, graft-versus-host disease, Alzheimer's disease, Parkinson's disease, asthma or fertility disorders. Furthermore, the polypeptides may be utilised as vaccines whilst the nucleic acids may be used as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New derivatives of human tropoelastin - with elastin-like vi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tropoelastin; hTE; elastin; glycosaminoglycan; GAG-binding; medical; pharmaceutical; veterinary; cosmetic application; anti-wrinkle; food; hand lotion; surgical implant; industrial product; human; SHEL; variant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human tropoelastin variant SHELdelta26A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fig 2; 82pp; English.
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100.0%; Pr
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RESULT 7
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        used where the wild-type protein would be degraded too easily, e.g. in contact with serum or wound exidate. The tropoclastin derivatives provide competitive inhibition of protease activity. The tropoclastin derivative derived protease-susceptibility sites, are useful in human or veterinary medicine, cosmetics (e.g. antiwrinkle or hand lotions), as bulking agents and for inducing chemotaxis. They are also useful for proliferation or growth inhibition, particularly of smooth muscle cells, epithelial or endothelial cells, fibroblasts, osteocytes, chondrocytes and plattelets. Peptidomimetics that mimic the protease cleavage site in tropoclastin derivatives are competitive inhibitors of the protease, and are used for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tropoelastin; derivative; SHEL-delta-26a; SHEL; proteolysis; protease; antiwrinkle; hand lotion; bulking agent; chemotaxis; proliferation; growth inhibition; peptidomimetic; lung damage; elastin; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                that susceptibility to proteolysis is reduced or eliminated, or a subsequence has been inserted so that susceptibility to proteolysis is increased. The derivatives have with reduced susceptibility, and can be
                                                                                                                                                                                                                                                     The present sequence represents a human reduced tropoelastin derivative, designated SHEL-delta-26a. The sequence is produced by removing exon 26a designated SHEL not defined). The protein is representative of tropoelastin derivatives of the invention, in which a subsequence has been mutated so
                                                                                                                                                                                                                                                                                                                                                                                                                             N-PSDB, AAZ61144.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    metastasis; blood clotting.
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                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 110-112; 136pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-182399/16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                          New derivatives of tropoelastin useful, e.g. for inducing chemotaxis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UNSY ) UNIV SYDNEY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence of a human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              698
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Pred. No. 3.9e-35;
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damage caused by

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RESULT 8
ADB40132
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Best Local S
Matches 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NOVX; cardiant; antiarteriosclerotic; hypotensive; cytostatic; anorectic; antidiabetic; immunosuppressive; anti-HIV; neuroprotective; nootropic; antiparkinsonian; antiasthmatic; yenticological; cardiomyopathy; atherosclerosis; hypertension; cancer; obesity; diabetes; AIDS; multiple sclerosis; graft-versus-host disease; Alzheimer's; Parkinson's; asthma; fertility disorder; vaccine; gene therapy; chromosome mapping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-AUG-2001;
03-AUG-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              controlling localized growth of cancers or metastases, or to limit protease activity that causes blood clotting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003064589-A2
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17-AUG-2001;
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                            CURAGEN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FPGALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLGVS
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2001US-0310951P
2001US-0311292P
2001US-0313201P
2001US-0313201P
2001US-0313415P
2001US-0313403P
2001US-0313702P
2001US-0314031P
2001US-0314031P
2001US-031466P
2001US-0315853P
2001US-0322716P
2001US-0322716P
2001US-0322716P
2001US-0323994P
2001US-03402339
2001US-03402339
2002US-036474P
2002US-0373825P
2002US-0373825P
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                              CORP
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100.0%;
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Pred. No.
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TEGENERAL X SERVICE SE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kekuda R, Miller CE, Patturajan M, Pena Shimkete RA, Zerhusen BD, Li L, Ji W, II Voss EZ, Boldog FL, Gorman L, Leite MW, Guo X, Zhong M, Gerlach VL, Hjalt T, Ræ Edinger SR, Ellerman K, Malyankar UM, Malsobrook JP, Lepley DM, Burgess CE, Maj
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                be utilised as vaccines whilst the nucleic acids may be used as hybridisation probes, in gene therapy, chromosome mapping, tissue preventive medicine and pharmacogenomics. The current sequence is the human NOV protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New NOVX polypeptides and nucleic acids, useful for preventing or treating NOVX-associated disorders, e.g. cancer, cardiomyopathy, atherosclerosis or diabetes, and in chromosome mapping, tissue typing
                                                                                                                                                                                                                                                                                                                         Tropoelastin; elastin; elastic fibre; smooth muscle cell proliferation; smooth muscle cell differentiation; smooth muscle cell migration; smooth muscle cell function; atherosclerosis; restenosis; aneurysm; vascular bypass graft stenosis; transplant arteriopathy; dissection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              invention demonstrates cardiant, antiarteriosclerotic, cytostatic, anorectic, antidiabetic, immunosuppressive,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N-PSDB;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a novel NOVX polypeptide. The invention demonstrates cardiant, antiarteriosclerotic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; SEQ ID NO 38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pharmacogenomics.
                          26-FEB-1999;
                                                                              28-FEB-2000;
                                                                                                                                        31-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Amino acid sequence of a human elastin polypeptide
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                                                                                                                                                                                                                                                                                                    hypertension; transplant arteriopathy.
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                                                                                    2000WO-US002526
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                             99US-00258217
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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, Padigaru M, ....
MW, Vernet CAM, Anders...
NH, Vernet CAM, Anders...
Stone DJ
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ches 0;
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Macdougall JR, Stone
Majumder K, Wolenc AR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
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RESULT 10
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AC AAW46
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 712 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Elastin based compositions useful for treating atherosclerosis, restenosis, vascular bypass graft stenosis, transplant arteriopathy, aneurysm, dissection SVAS and/or hypertension.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Keating MT,
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                                                                                                                        07-AUG-1996;
07-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human elastin containing non-natural polypeptide MFU-1 sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAW46315 standard; protein; 730 AA.
                                                                                                                                                                                                                      07-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                  WO9805685-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                             Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-JUL-1998 (first entry)
                                                                                                                                                                                                                                                                                    12-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MFU-1; minimal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            63
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                             PROTEIN SPECIALTIES LTD. HOSPITAL FOR SICK CHILDREN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FPGALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLGVS 124
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGVPGAIPGGVPGGVFYPGAGLGALGGGALGPGGKPLKPVPGGLAGAGLGAGLGAFPAVT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   coating;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   functional unit; elastin; human; fibrous protein;
ating; wound dressing.
                                                                                                                        96US-0023552P.
97US-00911364.
                                                                                                                                                                                                                      97WO-CA000560
                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers 374. .499
                                                                                                                                                                                                                                                                                                                                                                                                          /note= "MFU-1 polypeptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          79pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 515; DB 3;
Pred. No. 3.9e-35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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This represents the human elastin sequence containing the minimal CC functional unit (MFTU)-1 polypeptide of the invention. This MFTU-1 is a CC polypeptide that has at least 3 beta-sheet/ beta-turn structures, but is CC not a naturally occurring fibrous protein. Each beta-sheet structure has CC 1-2 (preferably 5-7) amino acids and the MFTU polypeptide may include at CC least 1 amino acid that can take part in crosslinking. The polypeptide CC can also be derived from the sequences of animal elastin, lamprin and CC spider silk protein. The MFTU polypeptides are self-aligning peptides CC spider silk protein. The MFTU polypeptides are self-aligning peptides of the ving the same primary structure as part of a natural fibrous protein. The MFTU polypeptides are self-aligning peptides or metal, particularly for use as blood vessel or heart valve replacements, CC wound or burn dressings, or stents. They can be used in cosmetic, elastic CC wound or burn dressings, or stents. They can be used in cosmetic, elastic CC prosthese based on the MFTU allow penetration of endothelial cells, so become permanent, living, tissue replacements. The MFTU polypeptides have better biocompatibility than known elastin-based materials. They are well cellials can be made from 2 or more different well polypeptides to allow properties to be tailored for particular applications, e.g. combining the high extensibility of elastin and the high tensile strength of spider can be made from 2 or more different well strength of spider well as the polypeptides to allow 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New non-natural polypeptide with multiple beta-sheet, beta-turn structures - particularly based on human elastin, useful for coprostheses, as wound dressings, etc., allows ingrowth of cells.
Sequence 730 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rothstein A,
                                                                  high extensibility silk protein
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                                                                                 3 GGVPGAIPGGVPGGVFYPGAGLGALGGGALGPGGKPLKPVPGGLAGAGLGAGLGAFPAVT
                                                                                                                           Similarity
                                                     GGVPGA1PGGVPGGVFYPGAGLGALGGGALGPGGKPLKPVPGGLAGAGLGAGLGAFPAVT
FPGALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLGVS 98
                          FPGALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLGVS 100
                                                                                                             98.3%;
ilarity 100.0%;
Conservative (
                                                                                                             %; Score 515; DB
%; Pred. No. 4e-
0; Mismatches
                                                                                                                          DB 2; Length 730; 4e-35;
                                                                                                              0
                                                                                                                Indels
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                                                          60
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RESULT 11
AAB08631
ID AAB08
AAB08631 standard;
                                             peptide;
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Fusion protein comprising human elastin and c-myc.

20-DEC-2000

(first entry)

Tropoelastin; elastin; elastic fibre; smooth muscle cell proliferation; smooth muscle cell differentiation; smooth muscle cell migration; smooth muscle cell function; atherosclerosis; restenosis; aneurysm; vascular bypass graft stenosis; transplant arteriopathy; dissection; SVAS; hypertension; transplant arteriopathy. Synthetic.

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28-FEB-2000; 2000WO-US002526
                                  31-AUG-2000
                                                                                                        Homo sapiens.
Unidentified.
                                                                     WO200050068-A2
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26-FEB-1999;

99US-00258217

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RESULT 12
AAO17360
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Best Local S
Matches 98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    proliferation, differentiation and migration in vivo. The elastin-based compositions comprise at least one elastic fibre, elastins, tropoelastins (or fragments of them) which have biological activities comprising: inhibiting the proliferation of smooth muscle cells in vivo; stimulating the differentiation of smooth muscle cell in vivo; and regulating the migration of smooth muscle cell in vivo; and regulating the migration of smooth muscle cells in vivo; and regulating the for the prophylaxis or treatment of a disorder characterized by diminished capacity to regulate smooth muscle cell function such as atherosclerosis, restenosis, vascular bypass graft stenosis, transplant arteriopathy, aneurysm and/or dissection. Disorders which may be treated
                                                                                                                                                                                   Human; endometriosis; DNA chip; fibronectin; p27; reticulocalbin; aldehyde dehydrogenase 6; gravin; phospholipase C epsilon; elastin; insulin-like growth factor binding protein-2; alpha-2 type IV collagen; transmembrane receptor PTK7; collagen type XVIII alpha 1; platelet derived growth factor receptor alpha; laminin M chain; subtilisin like protein PACE4; nidogen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence represents a fusion protein, comprising elastin and c-myc, preceded by a His tag. The protein is used compositions of the invention. The specification describes el compositions that are potent regulators of smooth muscle cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  restenosis, vascular bypass graft stenosis, transplant arteriopathy, aneurysm, dissection SVAS and/or hypertension.
                 25-SEP-2000; 2000DE-01048633
                                                 21-AUG-2001; 2001EP-00250300
                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                        Human elastin
                                                                                                                                                                                                                                                                                                                                          19-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                             AAO17360 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         also include SVAS (undefined), hypertension, and transplant arteriopathy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-533134/48
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                                                                                     27-MAR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3; Page 48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGVPGAIPGGVPGGVFYPGAGLGALGGGALGPGGKPLKPVPGGLAGAGLGAGLGAFPAVT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FPGALVPGGVADAAAYKAAKAGAGLGGVPGVGGLGVS 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGVPGAIPGGVPGGVPYPGAGLGALGGGALGPGGKPLKPVPGGLAGAGLGAGLGAFPAVT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         730 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FPGALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLGVS
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                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                          protein; 730
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 515; DB 3;
Pred. No. 4e-35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 730;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                describes elastin based
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ADQ19747
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AC ADQ19
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a method for the in vitro diagnosis of endometriosis by determining the amount of gene product from at least one specific gene in a patient sample and comparing this with the amount of gene product in a control sample. A reduced level is indicative of endometriosis. The gene products may be fibronectin, p27, reticulocalbin, aldehyde dehydrogenase 6, gravin, phospholipase C epsilon, elastin, insulin-like growth factor binding protein-2, alpha-2 type IV collagen, transmembrane receptor PTK7, collagen type XVIII alpha-1, platelet derived growth factor receptor alpha, laminin M chain, subtilisin like protein PACE4 or nidogen. The method is useful for initial diagnosis of endometriosis, and also for monitoring progress and treatment of the disease. The present sequence is human elastin
                           Early detection of soft tissue sarcoma comprises determining expression of a gene in a first soft tissue sample and a normal soft tissue sample and comparing the gene expression, also useful in treating soft tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 730 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               In vitro diagnosis and monitoring of endometriosis, comprises detecting reduced expression of specific gene products, e.g. from the fibronectin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hess-Stumpp H, Haend
Regidor P, Scotti S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human soft tissue sarcoma-upregulated protein - SEQ ID 2566
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADQ19747 standard; protein;
                                                                                                                                                                                                                                                                                                                         26-NOV-2002; 2002US-0429739P.
                                                                                                                                                                                                                                                                                                                                                                                   26-NOV-2003; 2003WO-US038193
                                                                                                                                                                                                                                                                                                                                                                                                                                             10-JUN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2004048938-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      soft tissue sarcoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-AUG-2004
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                                                                                                                                                                                                                                                                     (PROT-)
                                                                                                                                                2004-441208/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   87
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                                                                                                                                                                                                       Ginsburg
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FPGALVPGGVADAAAYKAAKAGAGLGGVPGVGGLGVS
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                                                                                                                                                                                                          ΨW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98.3%; but
100.0%; Pr
'''A 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cytostatic; gene therapy; vaccine; screening; human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              В,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 515; DB; Pred. No. 4e-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kraetzschmar J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kreft B,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 730;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Winterhager
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Example 2; SEQ ID

NO 2566;

210pp; English.

invention relates to

a novel method for detecting

g soft ti

tissue

sarcoma

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           맑
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           which comprises obtaining a first soft tissue sample from an individual and a normal soft tissue sample from the same or different individual, determining the expression of a gene in both samples and comparing the expression of the gene in both soft tissue samples and comparing the expression in the first soft tissue sample indicates the presence of soft tissue sarcoma. The method of the invention has cytostatic applications and may be useful for detecting soft tissue sarcoma, possibly via gene therapy or vaccine production. The nucleic acid sequences may be useful in diagnostic and screening applications. The current sequence is that of a human soft tissue sarcoma-upregulated protein of the invention. The current sequence is not shown within the specification per se but was submitted in CD format by the inventor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 730 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY69068 standard; protein; 731
                                                              WPI; 2000-182399/16.
N-PSDB; AAZ61146.
                                                                                                                                                                                                                                                                                                                                                                        peptidomimetic;
                                                                                                                                                                                                                                                                                                                                                                                                                           Amino acid sequence of a human tropoelastin splice form
                       cell growth.
                                   New derivatives of tropoelastin useful, e.g.
                                                                                                      Weiss AS
                                                                                                                                                         17-JUL-1998;
                                                                                                                                                                                   19-JUL-1999;
                                                                                                                                                                                                             27-JAN-2000.
                                                                                                                                                                                                                                        WO200004043-A1.
                                                                                                                                                                                                                                                                 Cleavage-site
                                                                                                                                                                                                                                                                              Cleavage-site
                                                                                                                                                                                                                                                                                           Cleavage-site
                                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                     Tropoelastin;
hand lotion; k
                                                                                                                                (UNSY ) UNIV
                                                                                                                                                                                                                                                                                                       leavage-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           87
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3 GGVPGAIPGGVPGGVFYPGAGLGALGGGALGPGGKPLKPVPGGLAGAGLGAGLGAFPAVT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FPGALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLGVS 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FPGALVPGGVADAAAYKAAKAGAGLGGVPGVGGLGVS 124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGVPGA1PGGVPGGVFYPGAGLGALGGGALGPGGKFLKPVPGGLAGAGLGAGLGAFPAVT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                     bulking
                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first
                                                                                                                                 SYDNEY.
                                                                                                                                                                                                                                                                                                                                                                                                 derivative;
                                                                                                                                                         98AU-00004723
                                                                                                                                                                                   99WO-AU000580
                                                                                                                                                                                                                                                                 441.
503.
515.
564.
                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98.3%;
100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      entry)
                                                                                                                                                                                                                                                                                                                                                                       tive; proteolysis; protease; antiwrinkle; agent; chemotaxis; proliferation; growth damage; elastin; cancer; metastasis; bloo
                                                                                                                                                                                                                                                                 .442
.504
.516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 515;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                          e; proliferation; growth inhibition;
cancer; metastasis; blood clotting.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 8;
4e-35;
                                      for inducing chemotaxis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 730;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         62
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Disclosure; Page 107-109; 136pp; English.

The present invention relates to a minimal functional unit (MFU) of human elastin polypeptide. This protein is useful in a cosmetic material or a

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RESULT 15
AAB66657
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CCCXSS PTTTX R X PX P X P X P X P X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X P X X 
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Best Local
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                                                                                                                                                                           Novel polypeptides that comprise three beta-sheet/beta-turn structures and are not naturally occurring fibrous protein, used to produce prosthesis suitable for implantation into humans, and cosmetic material
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Minimal function
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human elastin protein without signal peptide.
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                                                                                                   Claim 1; Fig 1; 39pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (PROT-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-JUN-2000; 2000WO-US017829
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200100666-A2
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                                                                                                                                                                                                                                                                                                                                           WPI; 2001-102886/11.
                                                                                                                                                                                                                                                                                                                                                                                                               Rothstein A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PROTEIN SPECIALTIES LTD HSC RES & DEV LP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FPGALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLGVS 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGVPGAIPGGVPGGVFYPGAGLGALGGGALGPGGKPLKPVPGGLAGAGLGAGLGAFPAVT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                               Keeley F, Rothstein S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   9908-00340736
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           unit; MFU; human; elastin prosthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein; 731 AA
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CC prosthetic material such as prosthesis for blood vessel replacements, for CC heart valve replacement, tissue replacement, for covering burns, for CC covering wounds and stents

XX

XX

XX

SQ Sequence 731 AA;

Query Match

Best Local Similarity 100.0%; Pred. No. 4e-35;

Matches 98; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Matches 98; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 3 GGVPGAIPGGVPGGVFYPGAGLGALGGGALGPGKPLKPVPGGLAGAGLGAFDAVT 62

Qy 63 FPGALVPGGVADAAAAYKAAKGAGLGGVPGVGGLGVS 100

Qy 63 FPGALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLGVS 100

Qy 63 FPGALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLGVS 98

Search completed: August 10, 2005, 07:33:50

Job time: 121.905 secs
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Result
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Maximum Match 100%
Listing first 45 su
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Maximum DB
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Sequence:
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     Query
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Copyright (c) 1993 - 2005 Compugen Ltd.
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US-09-340-736E-1
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US-08-678-039A-40
US-08-397-633A-78
US-08-212-237-4
US-08-212-237A-108
US-08-707-237A-108
US-08-707-237A-108
US-08-642-246-30
US-08-866-029-31
US-08-866-029-31
US-08-866-029-34
US-08-866-029-34
US-08-866-029-36
US-08-868-212-237-3
US-08-868-229-26
US-08-868-29-28
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Sequence 2, Appli Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 40, Appli Sequence 40, Appli Sequence 40, Appli Sequence 27, Appli Sequence 27, Appli Sequence 30, Appli Sequence 30, Appli Sequence 30, Appli Sequence 30, Appli Sequence 104, Appli Sequence 26, Appli Sequence 27, Appli Sequence 28, Appli Sequence 25, Appli Sequence 27, App
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28 171 32.6 262 1 US-08-397-633A-73 Sequence 73, Appl 168 32.1 696 3 US-08-806-029-25 Sequence 25, Appl 31 167.5 32.0 251 1 US-08-806-029-25 Sequence 25, Appl 32 166 31.7 1413 1 US-08-175-155-39 Sequence 27, Appl 33 166 31.7 1413 2 US-08-175-155-39 Sequence 45, Appl 34 166 31.7 1464 1 US-08-477-509B-74 Sequence 47, Appl 35 166 31.7 1464 3 US-08-477-509B-74 Sequence 74, Appl 36 166 31.7 1464 3 US-08-482-085B-74 Sequence 74, Appl 37 164 31.3 988 1 US-08-482-085B-74 Sequence 74, Appl 38 164 31.3 988 1 US-08-212-237-5 Sequence 28, Appl 39 164 31.3 988 3 US-08-212-237-5 Sequence 28, Appl 39 164 31.3 988 3 US-08-212-237-5 Sequence 27, Appl 39 164 31.3 988 3 US-08-212-237-5 Sequence 27, Appl 39 164 31.3 988 3 US-08-317-633A-75 Sequence 27, Appl 39 164 31.3 988 3 US-08-317-633A-75 Sequence 75, Appl 41 162 30.9 281 1 US-08-397-633A-75 Sequence 75, Appl 42 161.5 30.8 877 1 US-08-397-633A-64 Sequence 54, Appl 44 161.5 30.8 877 1 US-08-397-633A-64 Sequence 54, Appl 44 161.5 30.8 884 2 US-08-435-641-15 Sequence 54, Appl 54 161.5 30.8 884 2 US-08-435-641-15 Sequence 54, Appl 54 161.5 30.8 884 2 US-08-435-641-15
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ALIGNMENTS

## RESULT 1 US-08-464-700-2 Sequence 2, Application US/08464700 Patent No. 6232458 GENERAL INFORMATION: APPLICATION UNMBER: US/08/464,700 PILING DATE: 7-JUN-1995 CLASSIFICATION: 435 PRIOR APPLICATION: 435 PRIOR APPLICATION DATA: APPLICATION UNMBER: AU PL6520 APPLICATION UNMBER: AU PL6520 PILING DATE: 22-DEC-1992 PRIOR APPLICATION DATA: APPLICATION NUMBER: AU PL9661 PILING DATE: 28-JUN-1993 PRIOR APPLICATION DATA: APPLICATION UNMBER: PCT/AU93/00655 PILING DATE: 16-DEC-1993 APTORNEY/AGENT INFORMATION: NAME: Raiv Marry E INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: NAME: Bak, Mary E. REGISTRATION NUMBER: 31,215 REFERENCE/DOCKET NUMBER: GHC3 TELECOMMUNICATION INFORMATION: TELEPHONE: 215-540-9200 TELEFAX: 215-540-5818 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: LEM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS APPLICANT: WEISS, ANTHONY S APPLICANT: MARTIN, STEPHEN L TITLE OF INVENTION: SYNTHETIC POLYNUCLEOTIDES NUMBER OF SEQUENCES: 54 TYPE: amino acid TOPOLOGY: linear MOLECULE TYPE: prot SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA: CORRESPONDENCE ADDRESS: ADDRESSEE: Howson ar COUNTRY: ZIP: 194 STREET: Spring House CITY: Spring House STATE: Pennsylvania LENGTH: 19477 Spring House Corporate Cntr, PO 733 amino acids USA Howson and Howson GHC3USA Version

Query Match

100.0%;

Score 524;

DB 3;

Length 733;

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LENGTH: 730
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Patent No. (
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Patent No. 5969106
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Best Local (
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APPLICANT: KRAETZSCHWAR, JOERN
APPLICANT: KRAETZSCHWAR, JOERN
APPLICANT: KREFT, BERTHOLT
APPLICANT: WINTERHAGER, ELKE
APPLICANT: SCOTTI, SIMONE
APPLICANT: SCOTTI, SIMONE
TITLE OF INVENTION: METHOD FOR IN VITRO DIAGNOSIS OF ENDOMETRIOSIS
FILE REFERENCE: SCH-1789
CURRENT APPLICATION NUMBER: US/09/961,403
CURRENT APPLICATION NUMBER: US/09/961,403
CURRENT FILING DATE: 2001-09-25
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: ROTHSTEIN, Aser
APPLICANT: KEELY, Fred W.
APPLICANT: KEELY, Fred W.
APPLICANT: ROTHSTEIN, Steven J.
TITLE OF INVENTION: SELF-ALIGNING PEPTIDES MODELED ON HUMAN
TITLE OF INVENTION: BLASTIN AND OTHER FIBROUS PROTEINS
NUMBER OF SEQUENCES: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: HE-STUMPP,
APPLICANT: HAENDLER,
APPLICANT: KRAETZSCHW
APPLICANT: KREFT, BER
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: FOLEY & LARDNER
STREET: 3000 K Street, N.W.
                                                                                                                                                                           CITY: Washington
                                                                                                                                       COUNTRY:
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                      Version #1.30
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Best Local Similarity
Matches 98; Conserv
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                                                                                                                                                       Matches
                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: KEELEY, FRED
APPLICANT: ROTHSTEIN, STEVEN
APPLICANT: ROTHSTEIN, STEVEN
TITLE OF INVENTION: SELF-ALIGNING PEPTIDES MODELED ON HUMAN ELASTIN
TITLE OF INVENTION: AND OTHER FIBROUS PROTEINS
FILE REFERENCE: 041082/0110
CURRENT APPLICATION NUMBER: US/09/340,736E
CURRENT FILING DATE: 1997-08-29
PRIOR APPLICATION NUMBER: 08/911,364
PRIOR FILING DATE: 1997-08-07
PRIOR FILING DATE: 1997-08-07
                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/023,552
PRIOR FILING DATE: 1996-08-07
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.1
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: ROTHSTEIN,
                                                                                                                                                                                                                                                    LENGTH: 731
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 60 FILING DATE: 07-AUG-1996 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 530 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 04
TELECOMMUNICATION INFORMATION:
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TOPOLOGY: li
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Bent, Stephen A. REGISTRATION NUMBER: 2
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61
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                                                                                                   3 GGVPGAIPGGVPGGVFYPGAGLGALGGGALGPGGKPLKFVPGGLAGAGLGAGLGAGLGAFPAVT
                                                                                                                                                                            h 98.3%; Score 515; DB 4; Similarity 100.0%; Pred. No. 1.4e-37;
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FPGALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLGVS 98
                                                                          GGVPGAIPGGVPGGVFYPGAGLGALGGGALGPGGKPLKPVPGGLAGAGLGAGLGAFPAVT
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                                                                                                                                                           Mismatches
                                                                                                                                                                                              Length 731;
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US-08-678-039A-40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 40,
Patent No. 5
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Best Local :
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LENGTH: 731
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APPLICANT: HSC RESEARCH AND DEVELOPMENT LIMITED PARTNERSHIP
TITLE OF INVENTION: SELF-ALIGNING PEPTIDES MODELED ON HUMAN ELASTIN AND
TITLE OF INVENTION: OTHER FIBROUS PROTEINS
FILE REFERENCE: 041082/0112
CURRENT APPLICATION NUMBER: US/09/964,662
CURRENT FILING DATE: 2003-05-08
FRIOR APPLICATION NUMBER: 09/340,736
PRIOR FILING DATE: 1999-06-29
                                                                                MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Vers
SOFTWARE: PatentIn Release #1.0, Vers
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/678,039A
FILING DATE: 10-UUL-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Saxe, Stephen A.
REGISTRATION NUMBER: 38,609
REGISTRATION NUMBER: 38,609
REFERENCE/DOCKET NUMBER: 2323-120A
TELECOMMUNICATION INFORMATION:
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                     TELEFAX: 202-783-6031 INFORMATION FOR SEQ ID NO: 40:
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APPLICANT: Morris,
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENCES: 42
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ANDRESSEE: Rothwell, Figg, Ernst & Kurz, P.C.
ANDRESSEE: Thirteenth Street, N.W., Suite 701 East
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: U
ZIP: 20004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Washington
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5858662
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                                                                    202-624-1589
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Colleen A.

Diagnosis of Williams Syndrome and
Williams Syndrome Cognitive Profile by Analysis of
Presence or Absence of a LIM-Kinase Gene
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                                                                                                                                                                                                                                                                                               Version #1.30
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US-08-397-633A-78
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Best Local S
Matches 55
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APPLICANT: Cappello, Joseph
TITLE OF INVENTION: PRODUCTS COMPRISING SUBSTRATESCAPABLE
TITLE OF INVENTION: OF ENZYMATIC CROSS-LINKING
                                                                                                                                                                                                                                                                                                                       TELEFAX: (415) 398-324
TELEX: 910 277299
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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NAME: Rowland, Bertram I
REGISTRATION NUMBER: 20,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: A-58848-1/BIR PROP-011-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                           MOLECULE TYPE:
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CITY: San Francisco
                                                                                                                                                                                                                                                                    STRANDEDNESS: single
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                                                                                                                                                                                                                                                            TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
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                              132 GVGVPGGLKGGGGVPGVGV-PGVGVPGVGVPGAGAGSGAGAGSGAGAGSGAGAGSGVGVP 190
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                                                            39 LKPVPGGLAGAGLGAGLGAFPAVTFPGALVPG-GVADAAAYKAAKAGAGLG-----GVP 92
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Similarity 100.0%; F
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GVG--GLGV 99
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nilarity 42.6%;
Conservative
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b; Pred. No. 1.5e
0; Mismatches
                                                                                                                                                          Score 177.5; DB 1,
Pred. No. 3.7e-08;
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er, Suite 3400
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US-08-806-029-27
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                                                                                                                       Sequence 27, Application US/08806029 Patent No. 6380154
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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Best Local Similarity
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INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
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                                              APPLICANT: Cappello, Joseph
APPLICANT: Stedronsky, Erwin R.
TITLE OF INVENTION: Synthetic Proteins for in vivo Drug
TITLE OF INVENTION: Delivery and Tissue Augmentation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: A-TELECOMMUNICATION INFORMATION: TELEPHONE: 415-781-1989
                  NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Rowland, Bertram I
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APPLICATION NUMBER: US,
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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REGISTRATION NUMBER: 20,015
       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, Version #1.25
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                                                                                                                                                                                                                                                 125 GAGSVPGVGVPGVGV 139
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   832 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
Flehr, Hohbach, Test, Albritton & Herbert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 174; DB 1; Length 832; Pred. No. 1.3e-07;
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Suite 3400
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                                                                                                                                                                                                                                                                                                     Sequence 4, Application PC/TUS9502772
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (415) 398-3249
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                              COUNTRY: U.S.A.
ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: unknown MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 31,801
REFERENCE/DOCKET NUMBER: A-
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 11-MAR-1994
ATTORNEY/AGENT INFORMATION:
NAME: Trecartin, Richard F.
REGISTRATION NUMBER: 31,801
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 9
                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
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                                                                                                                                                                    STREET: FOUL _____
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APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
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                                                                                                                                                                                                     ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
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53; Conservative
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                                                                                                                                                                                                                                                                       Synthetic Proteins As Implantables
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               PCT/US95/02772
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Pred. No. 1.3e-07;
6; Mismatches 32
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Best Local Similarity
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                                                             PILING DATE: 29-DEC-19
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 22-APR-19
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Crissman, John W.
APPLICANT: Dorman, Mary A.
TITLE OF INVENTION: METHODS FOR PREPARING SYNTHETIC
TITLE OF INVENTION: REPETITIVE DNA
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 09-NOV-
                                                                                                                                                                                  CLASSIFICATION: 435
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Rowland, Bertram I
REGISTRATION NUMBER: 20,015
REFERENCE/DOCKET NUMBER: FP
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                                                                                                                                                APPLICATION NUMBER: US 01
FILING DATE: 29-DEC-1993
                                                                                                                                                                                                                     APPLICATION NUMBER: FILING DATE: 03-SE
                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
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Crissman, John W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ferrari,
                                                                                                                                                                                                                     03-SEP-1996
                                                                                                  22-APR-1993
                                                 OG-NOV-1990
 MBER: US 07/269,429
09-NOV-1988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 174; DB 5;
Pred. No. 1.3e-07;
6; Mismatches 32
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Suite 3400
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RESULT 12
US-08-642-246-30
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 936 amin-
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                                                                                                                                                                                 ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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APPLICATION NUMBER: 1
FILING DATE: 04-NOV-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/114,618
FILING DATE: 29-OCT-1987
                                                                                                                                SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/642,240
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                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: STEDRONSKY APPLICANT: CAPPELLO,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE:
REFERENCE/DOCKET NUMBER: A6
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: Tissue Adhesive Using Synthetic TITLE OF INVENTION: Crosslinking
                                             CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: ROWLAND, Bertram I
REGISTRATION NUMBER: 20015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Trecartin, Richard F. REGISTRATION NUMBER: 31,801
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TELEPAX: (415) 398-3249
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                                                                                                                   FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Application US/08642246
                                                                                                                                                                                                                                                                                                                     E: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT Four Embarcadero Center, Suite 200
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                                                                                                                                   US/08/642,246
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                                   A61127-1/BIR
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                                                                                                                                                                      Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; DB 2;
1.5e-07;
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RESULT 13
US-09-451-206-30
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                                      US-09-451-206-30
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Query Match
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                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 30:
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                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/642,246
FILING DATE: CURLOWND
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: STEDRONSKY, Erwin R.
CAPPELLO, Joseph
TITLE OF INVENTION: Tissue Adhesive Using Synthetic
                                                        MOLECULE TYPE: protein SEQUENCE DESCRIPTION: 8
                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: A61127-1/BIR TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    129 PGKGVPGVGVPGVGVPGVGVPGAGAGSGAGAGSGVGVPLGPLGPGVGVPGVGVPGVGVPG 188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          189 KGVPGVGVPGVGVPGVGVPGAGAGSGAGAGSGVGVPLGPLGPGVGVPGVGVPGVGV 244
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 GVPG-AIPG-GVPG----GVFYPGAGLGALGG------GALGPG------GK
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                                                                                                                                    TYPE: amino acids
                                                                                                                                                                                                                                                                                          NAME: ROWLAND, Bertram I REGISTRATION NUMBER: 20015
                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/09/451,206 FILING DATE: 29-No. 6423333-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT STREET: Four Embarcadero Center, Suite 200 CITY: San Francisco
                                                                                                                STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: <Unknown>
                                                                                              TOPOLOGY: linear
                                                                                                                                                                                                                TELEFAX: 415-398-3249
                                                                                                                                                                                                                                    TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 936 amino acids
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ilarity 34.1%;
Conservative
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                                                          SEQ ID NO:
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Pred. No. 1.5e-07;
Score 174;
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                                                          30:
DB 4;
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Length 936;
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PCT-US96-06229-30
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                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
ATMAGE: ROWLAND, Bertram I.
REGISTRATION NUMBER: 20015
REFERENCE/DOCKET NUMBER: A611
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA: APPLICATION NUMBER: PCT/US96/0622 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Tissue Adhesi
TITLE OF INVENTION: Crosslinking
NUMBER OF SEQUENCES: 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
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189 KGVPGVGVPGVGVPGVGVPGAGAGSGAGAGSGVGVPLGPLGPGVGVPGVGVPGVGV 244
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                                 58 --FPAVTFPGALVPGGVADAAAAYKAAKAGAGLG------
                                                                                                          38 PLKPVPG-----GLAGAGLGAGLGA-----
                                                                                                                                              69 GVPGKGVPGVGVPGVGVPGAGAGSGAGAGSGVGVPLGPLGPGVGVPGVGVPGVGV 128
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                                                                       PGKGVPGVGVPGVGVPGVGVPGAGAGSGAGAGSGVGVPLGPLGPGVGVPGVGVPGVGVPG 188
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CAPPELLO, Joseph
NVENTION: Tissue Adhesive Using Synthetic
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                                                                                                                                                                                                                                      Score 174; DB 5;
Pred. No. 1.5e-07;
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7; Mismatches 29
                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                   ---GALGPG-----GK 37
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                                     -GVPGVG--GLGV 99
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Search completed: August 10, 2005, 07:39:40 Job time: 30.8507 secs
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US-08-806-029-34
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTION DATA:
APPLICATION UNMBER: US/08/806,029
FILING DATE: 24-FEB-1997
CLASSIFICATION NUMBER: US 08/212,237
APPLICATION NUMBER: US 08/212,237
ATTORNEY/AGENT INFORMATION:
REGERENCE/DOCKET NUMBER: A-58847-2/RFT/MTK
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 34, Application US/08806029 Patent No. 6380154
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APPLICANT: Cappello, Joseph
APPLICANT: Stedronsky, Erwin R.
TITLE OF INVENTION: Synthetic Proteins for in vivo Drug
TITLE OF INVENTION: Delivery and Tissue Augmentation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (415) 781-1989
TELEPAX: (415) 398-3249
INFORMATION FOR SEQ ID NO: 34:
SEQUENCE CHARACTERISTICS:
LENGTH: 312 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS: Fehr, Hohbach, Test,
ADDRESSEE: Flehr, Hohbach, Test,
STREET: Four Embarcadero Center,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: unkr
TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: San Francisco
STATE: California
                                                                                                      150 VPGVGVPGAGAGSGAGAGSGVGVPGVGVPGVGVPGVGV 187
                                                                                                                                                         68 VPGGVADAAAAYKAAKAGAGLG----GVPGVG--GLGV 99
                                                                                                                                                                                                            30
                                                                                                                                                                                                                                                            44 G-----FPAVTFPGAL 67
                                                                                                                                                                                                                                                                                                                                                             4 GVPG-AIPG-GVPG----GVFYPGAGLGALGGGALGPG------GKPLKPVP 43
                                                                                                                                                                                                                                                                                                               United States
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                                                                                                                                                                                                                                                                                                                                                                                                             7; Mismatches
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Suite 3400
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OM protein - protein search, using sw model
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Maximum Match 100%
Listing first 45 s
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Maximum DB seq length: 200000000
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Perfect score:
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                 Published Applications AA:*

1: /cgn2_6/ptodata/1/pubpaa/US07 PUBCOMB.pep:*
2: /cgn2_6/ptodata/1/pubpaa/US06 NEW PUB.pep:*
3: /cgn2_6/ptodata/1/pubpaa/US06 NEW PUB.pep:*
4: /cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
5: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
6: /cgn2_6/ptodata/1/pubpaa/US08_NEW PUB.pep:*
7: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
8: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
9: /cgn2_6/ptodata/1/pubpaa/US09A_PUBCOMB.pep:*
10: /cgn2_6/ptodata/1/pubpaa/US09A_PUBCOMB.pep:*
11: /cgn2_6/ptodata/1/pubpaa/US09C_PUBCOMB.pep:*
13: /cgn2_6/ptodata/1/pubpaa/US09C_PUBCOMB.pep:*
13: /cgn2_6/ptodata/1/pubpaa/US09C_PUBCOMB.pep:*
14: /cgn2_6/ptodata/1/pubpaa/US09C_PUBCOMB.pep:*
15: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*
16: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
17: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
18: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
19: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
19: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
19: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
19: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
19: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
19: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
10: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
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524
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Copyright (c) 1993 - 2005 Compugen Ltd.
6/ptodata/1/pubpaa/US60_PUBCOMB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

17 174	10 174	9 334.5	8 448	7 515	6 515	5 515	4 515	3 515	2 515	1 515	Result No. Score
			85.5								Query
936	832	663	617	788	788	731	730	730	711	692	Query Match Length DB
14	œ	15	15	16	16	10	16	10	15	15	
US-10-117-931-30	US-08-806-029-27	US-10-108-260A-2477	5 US-10-104-047-2915	US-10-852-575-3	US-10-852-065-4	US-09-964-662-1	US-10-723-860-2566	US-09-961-403-8	US-10-210-172-38	US-10-210-172-40	ID
Sequence 30, Appl	Sequence 27, Appl	Sequence 2477, Ap	Sequence 2915, Ap	Sequence 3, Appli	Sequence 4, Appli	Sequence 1, Appli	Sequence 2566, Ap	Sequence 8, Appli	Sequence 38, Appl	Sequence 40, Appl	Description

## ALIGNMENTS

US-10-210-172-40 GENERAL Sequence 40, Application US/10210172 Publication No. US20040043928A1 APPLICANT:
APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT: INFORMATION: Boldog, Ferenc Gorman, Linda Leite, Mario Vernet, Corine Casman, scur Malyankar, Uriel MacDougall, John Stone, David Hjalt, Tor Rastelli, Patturajan, Meera Pena, Carol Rieger, Daniel Miller, Charles Rastelli, Luca Spytek, Kimberly Edinger, Shlomit Zhong, Mei Gerlach, Valerie Anderson, David Guo, Xiaojia Padigaru, Muralidhara Casman, Stacie Shimkets, Richard Zerhusen, Bryan Ellerman, Karen Weizhen Tord Ramesh John

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US-10-210-172-38
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; ORGANISM: Homo sapiens
US-10-210-172-40
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Best Local S
Matches 98
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LENGTH: 692
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PRIOR FILING DATE: 2001-08-02
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APPLICATION NUMBER: 60/373,814
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OF INVENTION: THERAPEUTIC POLYPEPTIDES,
REFERENCE: 21402-416 A
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                                                                                                                         Casman, UC
Toss, Edward
                                                                                                                                         Padigaru, Muz.
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Zhong, Mei
Gerlach, Val
Hjalt, Tord
                                                                                       Boldog, Ferenc
Gorman, Linda
Leite, Mario
                                                                                                                                                                                                          Rieger, Daniel
Shimkets, Richard
Zerhusen, Bryan
                                            Guo, Xiaojia
                                                          Vernet, Corine
Anderson, Davi
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Pena, Carol
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/ 100.0%; Pr
                                                                                                                                                                                                                                                                                         Charles
                                                          David
                                                                                                                                                                  Muralidhara
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Sequence 8, Application US/09961403
Publication No. US20030077589A1
GENERAL INFORMATION:
APPLICANT: HE-STUMPP, HOLGER
APPLICANT: HAENDLER, BERNARD
APPLICANT: KRAETZSCHMAR, JOERN
APPLICANT: KRAETZSCHMAR, JOERN
APPLICANT: KRAETZSCHMAR, JOERN
APPLICANT: KRAETZSCHMAR, JOERN
APPLICANT: MINTERHAGER, ELKE
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US-09-961-403-8
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LENGTH: 711
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TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES,
FILE REPERBOCE: 21402-416 A
CURRENT APPLICATION NUMBER: US/10/210,172
CURRENT FILING DATE: 2001-08-01
PRIOR APPLICATION NUMBER: 60/309,501
PRIOR FILING DATE: 2001-08-02
PRIOR APPLICATION NUMBER: 60/323,994
PRIOR PRI
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PRIOR
                            APPLICANT: REGIOOR, PEDRO
APPLICANT: SCOTTI, SIMONE
TITLE OF INVENTION: METHOD FOR IN VITRO DIAGNOSIS
FILE REPERENCE: SCH-1789
FULRANT APPLICATION NUMBER: US/09/961,403
CURRENT FILLING DATE: 2001-09-25
NUMBER OF SEQ ID NOS: 15
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PRIOR APPLICATION NUMBER: 60/312,892
PRIOR FILING DATE: 2001-08-16
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PRIOR FILING DATE: 2001-08-03
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PRIOR APPLICATION NUMBER: 60/373,814
PRIOR FILING DATE: 2002-04-19
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SOFTWARE: PatentIn Ver.
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ORGANISM: Homo sapiens
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APPLICATION NUMBER: 60/311,979
FILING DATE: 2001-08-13
APPLICATION NUMBER: 60/313,201
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KREFT, BERTHOLT
WINTERHAGER, ELKE
REGIDOR, PEDRO
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Malyankar, Uriel
MacDougall, John
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Spytek, Kimberly
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US-10-723-860-2566
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APPLICANT: Cinsburg, Wendy M.
APPLICANT: Cinsburg, Wendy M.
APPLICANT: Zlotnik, Albert
TITLE OF INVENTION: Methods of Diagnosis of Soft Tissue Sarcoma, Compositions of Invention of Invention of Invention of Screening for Soft Tissue Sarcoma Modulators
FILE REFERENCE: 05802.0193.NPUS01
CURRENT APPLICATION NUMBER: US/10/723,860
CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: 60/429,739
PRIOR APPLICATION NUMBER: 60/429,739
PRIOR FILING DATE: 2002-11-26
NUMBER OF SEQ ID NOS: 8393
SOFTWARE: Patentin version 3.2
SEQ ID NO 2566
LENGTH: 730
         APPLICANT: PROTEIN SPECIALTIES LID.
APPLICANT: PROTEIN SPECIALTIES LID.
APPLICANT: HSC RESEARCH AND DEVELOPMENT LIMITED PARTNERSHIP
APPLICANT: HSC RESEARCH AND DEVELOPMENT LIMITED PARTNERSHIP
TITLE OF INVENTION: SELF-ALIGNING PEPTIDES MODELED ON HUMAN ELASTIN AND
TITLE OF INVENTION: OTHER FIBROUS PROTEINS
FILE REFERENCE: 041082/0112
CURRENT PILING DATE: 2003-05-08
PRIOR APPLICATION NUMBER: 09/340,736
PRIOR APPLICATION NUMBER: 09/340,736
PRIOR APPLICATION NUMBER: 09/340,736
PRIOR PILING DATE: 1099-06-29
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.1
SOFTWARE: Patentin Ver. 2.1
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LENGTH: 730
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Best Local Similarity
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Best Local Similarity
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
LENGTH: 731
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Sequence 3, Application US/10852575
Publication No. US20040258676A1
GENERAL INFORMATION:
APPLICANT: PERRIER, Eric
APPLICANT: BOUEZ, Charbel
APPLICANT: CENIZO, Valerie
APPLICANT: CENIZO, Valerie
APPLICANT: DAMOUR, POSCAI
APPLICANT: GLEYZAL, Claudine
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US-10-852-065-4
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; LENGTH: 788
; TYPE: PRT
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Best Local Similarity 100.0%;
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APPLICANT: DAMOUR, Odile
APPLICANT: GLEYZAL, Claudine
APPLICANT: ANDRE, Val, rie
APPLICANT: ANDRE, Val, rie
APPLICANT: REYMERNIER, Corinne
TITLE OF INVENTION: Stimulation of the synthesis and of the activity of an isoform
TITLE OF INVENTION: of 19syl oxidase-like LOXL for stimulating the formation of
TITLE OF INVENTION: elastic fibers
TITLE OF INVENTION: elastic fibers
TITLE OF INVENTION NUMBER: US/10/852,065
CURRENT FILING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: FR 03 07177
PRIOR FILING DATE: 2003-06-13
NUMBER OF SEQ ID NOS: 31
NUMBER OF SEQ ID NOS: 31
NUMBER OF SEQ ID NOS: 31
NUMBER: Patentin version 3.1
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ORGANISM:
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DAMOUR, Odile
GLEYZAL, Claudine
ANDRE, Valerie
REYMERMIER, Corinne
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CENIZO, Val, rie
BOUEZ, Charbel
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  Corinne
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                                                                                                                                                                                        US-10-108-260A-2477
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; ORGANISM: Homo sapiens US-10-104-047-2915
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Best Local S
Matches 98
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SOFTWARE: PatentIn Ver. 2.1
SEQ_ID_NO 2915
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 2915, Application US/10104047 Publication No. US20030236392A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 3
LENGTH: 788
                                                                                                                         Sequence 2477, Application US/10108260A Publication No. US20040005560A1 GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: HELIX RESEARCH INSTITUTE
TITLE OF INVENTION: NO. US20030236392A1el full length cDNA
FILE REFERENCE: H1-A0105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT FILING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: FR 03 07178
PRIOR FILING DATE: 2003-06-13
NUMBER OF SEQ ID NOS: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Stimulation of the activity of an isoform of lysyl oxidase TITLE OF INVENTION: for combating against some pathologies due to an incomplete, TITLE OF INVENTION: absent or disorganized elastogenesis FILE REFERENCE: 11123,82US01 CURRENT APPLICATION NUMBER: US/10/852,575
CURRENT FILING DATE: 2004-05-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/104,047
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER:
PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn version 3.1
FILE REFERENCE: H1-A0106
CURRENT APPLICATION NUMBER: US/10/108,260A
CURRENT FILING DATE: 2002-03-27
NUMBER OF SEQ ID NOS: 5458
                                                                               APPLICANT: HELIX RESEARCH INSTITUTE
TITLE OF INVENTION: No. US20040005560Alel full length cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                              FPGALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLGVS 114
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Pred. No. 2.2e-26;
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RESULT 10
US-08-806-029-27
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                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (415) 398-3249
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
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APPLICANT: Stedronsky, Erwin R.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/806,029
FILING DATE: 24-FEB-1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/212,237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PATENTIA Release #1.0.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 31,801
REFERENCE/DOCKET NUMBER: TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 11-MAR-1994
ATTORNEY/AGENT INFORMATION:
NAME: Trecartin, Richard F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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   10 GVPGVGVPGVGVPGVGVPGVGVPGVGVPGVGCAGAGSGAG-----AGSGAGAGSGAGAGS
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                                                          4 GVPG-AIPG-GVPG----GVFYPGAGLGALGGGALGPGGKPLKPVPGGLAGAGLGAGLGA
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                                                                                                                                                                      Score 174; DB 8;
Pred. No. 2.3e-05;
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TOPOLOGY: linear
; MOLECULE TYPE: protein
; SEQUENCE DESCRIPTION: SEQ ID NO:
US-10-117-931-30
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US-10-117-931-30
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RESULT 12
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Publication No. US20030104589A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 415-398-3249 INFORMATION FOR SEQ ID NO: 30:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: ROWLAND, Bertram I
REGISTRATION NUMBER: 20015
REFERENCE/DOCKET NUMBER: A61127-1/BIR
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/642,246
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/117,931
FILING DATE: 05-Apr-2002
CLASSIFICATION: <Unknown>
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT STREET: Four Embarcadero Center, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CAPPELLO, Joseph
TITLE OF INVENTION: Tissue Adhesive Using Synthetic
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                                                                                  --FPAVTFPGALVPGGVADAAAAYKAAKAGAGLG------GVPGVG--GLGV 99
                                                                                                                  PGKGVPGVGVPGVGVPGVGVPGAGAGSGAGAGSGVGVPLGPLGPGVGVPGVGVPGVGVPG 188
                                                                                                                                                      PLKPVPG---
                                                                                                                                                                                      GVPGKGVPGVGVPGVGVPGVGVPGAGAGAGSGAGAGSGVGVPLGPLGPGVGVPGVGVPGVGV 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: San Francisco
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GLGGVPGVG--GLGV 99
                                                  KGVPGVGVPGVGVPGVGAGAGAGSGAGAGSGVGVPLGPLGPGVGVPGVGVPGVGV 244
                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 936 amino acids TYPE: amino acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: US
                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  94111
                                                                                                                                                                                                                                                                        33.2%;
34.1%;
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                                                                                                                                                                                                                        -GVFYPGAGLGALGG------
                                                                                                                                                      GLAGAGLGAGLGA----
                                                                                                                                                                                                                                                      Score 174; DB 14;
Pred. No. 2.5e-05;
7; Mismatches 29;
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                                                                                                                                                                                                                                                                                        Length 936;
                                                                                                                                                                                                                          GALGPG-----GK 37
                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                        80;
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Best Local :
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RESULT 13
US-10-117-931-26
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publication No. US20020045567A1.

GENERAL INFORMATION:
APPLICANT: Cappello, Joseph
APPLICANT: Stedronsky, Erwin R.
APPLICANT: Stedronsky, Erwin R.
TITLE OF INVENTION: Delivery and Tissue Augmen
                                                                                               Sequence 26, Application US/10117931
Publication No. US20030104589A1
GENERAL INFORMATION:
APPLICANT: STEDRONSKY, Erwin R.
CAPPELLO, Joseph
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (415) 398-324
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Trecartin, Richard F.
REGISTRATION NUMBER: 31,801
REFERENCE/DOCKET NUMBER: A-58847-2/RFT/MTK
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/806,029 FILLING DATE: 24-FEB-1997 CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 312 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 11-MAR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: un
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: San Francisco
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CORRESPONDENCE ADDRESS:
REST, ALBRITTON & HERBERT
                                                              TITLE OF INVENTION: Tissue Adhesive Using Crosslinking
                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                   44 G-----GLAGAGLGAGLGA-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 GVPG-AIPG-GVPG----GVFYPGAGLGALGGGALGPG------
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                 VPGVGVPGAGAGSGAGAGSGVGVPGVGVPGVGVPGVGV 187
                                                                                                                                                                                                                                                                                                      VPGGVADAAAAYKAAKAGAGLG----GVPGVG---GLGV
                                                                                                                                                                                                                                                                                                                                             amino acid
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36.1%;
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Center,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 173; DB 8;
Pred. No. 1.1e-05;
7; Mismatches 32;
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Suite 3400
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Best Local S
Matches 57
                                                                                                                                                                                                                                                                                                                                                   Sequence 26, App
Publication No.
                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO: 26:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                            APPLICANT: Cappello, Joseph
APPLICANT: Stedronsky, Erwin R.
TITLE OF INVENTION: Synthetic Proteins for in vivo Drug
TITLE OF INVENTION: Delivery and Tissue Augmentation
                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
                                                                                                                                                                       NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
                                                                                                                 COUNTRY:
                                                                                                                                                       CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: ROWLAND, Bertram I
REGISTRATION NUMBER: 20015
REFERENCE/DOCKET NUMBER: A61127-1/BIR
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/117,931
FILING DATE: 05-Apr-2002
CLASSIFICATION: <Unknown>
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/642,246
FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: protein SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       183 VPGVGVPGAGAGSGAGAGSGVGVPGVGVPGVGVPGVGV 220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 68 VPGGVADAAAYKAAKAGAGLG----GVPGVG--GLGV 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        44 G-----GLAGAGLGAGLGA-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ch 33.0%; 1 Similarity 36.1%; 57; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4 GVPG-AIPG-GVPG----GVFYPGAGLGALGGGALGPG-----GKPLKPVP
                                                                                                 94111
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STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 378 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
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                                                                                                                                     California
                                                                                                                                                                                                                                                                                                                                                     Application US/08806029
                                                                                                                   United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 173; DB 14; Length 378; Pred. No. 1.3e-05;
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  Version #1.30
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RESULT 15
US-10-117-931-25
; Sequence 25, Application US/10117931
; Publication No. US20030104589A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TOPOLOGY: unknown ; MOLECULE TYPE: protein US-08-806-029-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 34.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (415) 398-3249 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 11-MAR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Trecartin, Richard REGISTRATION NUMBER: 31,8 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 08/212,237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CENGTH:
                                                                                                                             COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/117,931
FILING DATE: 05-Apr-2002
CLASSIFICATION: <Unknown>
                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/642,246
FILING DATE: <UNKNOWN>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT STREET: Four Embarcadero Center, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CAPPELLO, Joseph TITLE OF INVENTION: Tissue Adhesive Using Synthetic
                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: STEDRONSKY, Erwin R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   130 VPGVGVPGVGVPGVGGAGAGSGAGAGSVPGVGVPGVGV 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          71 -----GVADAAAAYKAAKAGAGLGGVPGVG--GLGV 99
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                                                                                                                                                                                                                                                                                                                                                   STATE: CA
                NAME: ROWLAND, Bertram I REGISTRATION NUMBER: 200
                                                                                                                                                                                                                                                                                                             ZIP: 94111
                                                                                                                                                                                                                                                                                                                                                                  CITY: San Francisco
                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                             COUNTRY: US
REFERENCE/DOCKET NUMBER: A61127-1/BIR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                936 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Crosslinking
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Pred. No. 3e-05;
*** matches 35;
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
TELEPHAX: 415-781-1989
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 1002 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 25:
US-10-117-931-25
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Search completed: August 10, 2005, 07:56:34
Job time : 105.478 secs
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                                                              44 G-----GLAGAGLGAGLGA-----
                                                                                                                                                                                                                 4 GVPG-AIPG-GVPG----GVFYPGAGLGALGGGALGPG-------GKPLKPVP 43
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Result
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Maximum Match 100%
Listing first 45 s
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Title:
Perfect score:
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       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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m24782; NID:g182063; PIDM:AAA53190. poelastin refers to a soluble precu [:119107; OMIM:130160 -7q11.23 equence #status predicted <8IG> in #status predicted <mat> ds: #status predicted ds: #status predicted</mat>	2061; PIDN:AAA5238 site of several sp H.; Chu, M.L.; Dav of human elastin c B; PMID:2831431	;182052; PIDN:AAA52379. A.; Baldwin, C.T.; Ind 188 ttin cDNAs from a human 109960; PMID:3171221	Sheppard, I 987 indic D:3039501  UMK5; GB:M:oldstein, r oldstein; r n elastin c D:2722804	96 #text	30 125 23.9 1147 1 MWAXIB 31 124.5 23.8 307 2 T27609 32 124.5 23.8 749 2 A70812 33 124 23.7 228 2 T49891 34 124 23.7 515 2 H70663 35 124 23.7 515 2 H70663 36 124 23.7 1660 2 A70869 37 123.5 23.6 479 2 F70573 38 123.5 23.6 512 2 S70644 39 123.5 23.6 512 2 T00799 40 123 23.5 256 2 A70514 41 122.5 23.4 271 2 S34666 42 122.5 23.4 271 2 S34666 43 121.5 23.2 112 2 JQ1064 44 121.5 23.2 173 2 JQ1064 45 121.5 23.2 220 2 T14441
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R; Franzblau, C.;

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G.D.; Mogayzel

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C;Superfamily: elastin
C;Keywords: alternative splicing; extracellular matrix;
F;1-27/Domain: signal sequence #status predicted <SIG>
F;28-860/Product: elastin #status predicted <MAT>
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A;Title: Use of an intron length polymorphism to localize the A;Reference number: A55721; MUID:95130069; PMID:7829060
               A;Cross-references: UNIPROT:099372; GB:M60647; GB:J05292; NID:g207444; PIDN:AAA42269.1; R;Deak, S.B.; Pierce, R.A.; Belsky, S.A.; Riley, D.J.; Boyd, C.D.
J. Biol. Chem. 263, 13504-13507, 1988
A;Title: Rat tropoelastin is synthesized from a 3.5-kilobase mRNA.
A;Reference number: A30878; NUID:88330868; PMID:2971041
A;Accession: A30878; NUID:88330868; PMID:2971041
A;Accession: A30878
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 781-864 <DEA>
A;Residues: 781-864 <DEA>
                                                                                                                                                                                                                                                                                                                            N;Alternate names: tropoelastin
C;Species: Rattus norvegicus (Norway rat)
C;Date: 11-Jan-1991 #sequence_revision 16-Aug-1996 #text_change 09-Jul-2004
C;Accession: A36106; A30878; A36523; S02173; I54172; I68505
R;Pierce, R.A.; Deak, S.B.; Stolle, C.A.; Boyd, C.D.
Biochemistry 29, 9677-9683, 1990
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C;Date: 18-Aug-1995 #sequence_revision 16-Aug-1996 #text_change
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A; Residues: 1-860 < WYD>
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GB:J04035; NID:g207442; PIDN:AAA42268.1; PID:g207443
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Pred. No. 1.4e-30;
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RESULT 4
S59623
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C;Keywords: alternative splicing; extracellular matrix;
F;1-21/Domain: signal sequence #status predicted <SIG>
F;22-864/Product: elastin #status predicted <MAT>
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A;Status: translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 558-864 <RE2>
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A, Residues: 264-533
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A;Title: Elements of the rat troppelastin gene associated A;Reference number: I54172; MUID:92241859; PMID:1572637
A;Accession: I54172
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A;Title: Characterization of rat heart tropoelastin.
A;Reference number: S02173; MUID:89117149; PMID:2913
A;Accession: S02173
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A; Accession: A36523
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A; Title: Role of tropoelastin
                                                      A;Cross-references: UNIPROT:P11547
R;Yoon, K.; Davidson, J.M.; Boyd, C.;
Arch. Biochem. Biophys. 241, 684-691,
A;Title: Analysis of the 3' region of
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A; Residues: 'IP', 369-545, 548-764, 770-864
                  A; Reference number: A24758; A; Accession: A24758
                                                                                                                                          A; Molecule type: mRNA
A; Residues: 1-770 < MAU>
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A; Status: preliminary;
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type: mRNA
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264, 15115-15119, 1989
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; PMID:3839997
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R;Rosenbloom, J.

Lab. Invest. 51, 605-623, 1984
A;Title: Biology of disease: Elastin: Relation of protein and gene structure to A;Reference number: I45885; MUID:85059254; PMID:6150137.
A;Accession: I45886
A;Status: preliminary; translated from GB/FMPT'/~--
A;Molecule type: DNA
A;Residues: 679-677
A; Molecule type: DNA
A; Residues: 678-683,685-747 <ROS>
A; Cross-references: GB:M31898, NID:9163015; PIDN:AAA96417.1;
A; Cross-references: GB:M31898, NID:9163015; PIDN:AAA96417.1;
A; Fishem. Biophys. Res. Commun. 186, 549-555, 1992
A; Title: The cysteine residues in the carboxy terminal domain A; Reference number: A58621; MUID:92337651; PMID:1632791
A; Contents: annotation, disulfide bonds
C; Comment: The term type:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  procupentstry 24, 3075-3080, 1985
A;Title: Structure of the 3' portion of the bovine elastin gene A;Recession: A22343; MUID:85280426; PMID:2992576
A;Molecula ***
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R;Raju, K.; Anwar, R.A.
J. Biol. Chem. 262, 5755-5762, 1987
A;Title: Primary structures of bovine elastin a, b, and c ded A;Reference number: A92640; MUID:87194772; PMID:3032943
A;Accession: A26728
A;Molecule type: mRNA
A;Residues: 1,'RS',4-11,'E',13-636,'V',638-747 <RAJ>
A;Cross-references: GB:J02717; NID:g163019; PIDN:AAA30503.1;
A;Cross-references: GB:J02717; NID:g163019; PIDN:AAA30503.1;
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A;Residues: 1,'RS',4-11,'E',13-225,260-636,'V',638-747 <RA3>
A;Residues: 1,'RS',4-11,'E',13-225,260-636,'V',638-747 <RA3>
A;Cross-references: GB:K03506; NID:g163027; PIDN:AAA30506.1; PID:g163028
R;Cicila, G.; May, M.; Ornstein-Goldstein, N.; Indik, Z.; Morrow, S.; Yel
Biochmistry 24, 3075-380, 1985
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A;Accession: B26728
A;Molecule type: mRNA
A;Residues: 1,'RS',4-11,'E',13-225,240-636,'V',638-747 <RA2>
A;Cross-references: GB:K03505; NID:g163025; PIDN:AAA30505.1;
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C;Superfamily: elastin
C;Keywords: alternative splicing; extracellular matrix; glycoprotein;
F;760-765/Disulfide bonds: #status predicted
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A; Residues: 1-27 < YEH>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AFPGGFFGAGGGAAGAAAYKAAAKAGAGGLGGVGGIGGVGGLGVS
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Pred. No. 8.1e-17;
     refers to a soluble precursor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4.
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     of the extracellular
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--AGLGAGLGAFPAVTFPGALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLG

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C;Superfamily: elastin
C;Keywords: alternative splicing; extracellular matrix; glycoprotein; hydroxylysine
F;1-747/Product: elastin precursor, splice form a #status predicted <EPA>
F;1-225,260-747/Product: elastin precursor, splice form c #status predicted <EPC>
F;1-225,240-747/Product: elastin precursor, splice form b #status predicted <EPB>
F;1-26/Domain: signal sequence #status predicted <SIG>
F;1-26/Domain: elastin #status predicted <MATP
F;27-747/Product: elastin #status predicted <MATP
F;105,09,252,271,275,324,337,400,404,407,445,448,489,493,544,548,552,606,609,645,649,F;737-742/Disulfide bonds: #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Nolecule type: mRNA
A;Nolecule type: mRNA
A;Residues: 85-784 <BAU>
A;Cross-references: GB.M21880; NID:g212741; PIDN:AAA49082.1; PID:g212742
A;Cross-references: GB.M21880; Nibhikawa, T.; Tajima, M.; Fukasawa, T.
                                                                                                                                                                                                                       C;Keywords: alternative splicing; extracellular matrix; glycoprotein; F;1-24/Domain: signal sequence (fragment) #status predicted <SIG>F;25-212,237-524,535-784/Product: elastin #status predicted <MAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                  R;Tokimitsu, I.; Tajima, S.; Nishikawa, T.;
Arch. Biochem. Biophys. 256, 455-461, 1987
A;Tille: Sequence analysis of elastin cDNA
A;Reference number: A27264; MUID:87297534;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R;Baule, V.J.; Foster, J.A.
Biochem. Biophys. Res. Commun. 154, 1054-1060, 1988
Biochem. Nultiple chick tropoelastin mRNAs.
A;Reference number: A30795; MUID:88309083; PMID:284:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Molecule type: mRNA
A;Residues: 1-212,237-524,535-784 <BR2>
A;Cross-references: UNIPROT:P07916; GB:M15889; NID:g212803;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Species: Gallus gallus (chicken)
C;Date: 05-Oct-1988 #sequence revision 26-Jul-1996
C;Accession: A26601; A30795; A27264
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A;Residues: 491-569,'G',571-604,'A',606-643,'A',645-687,'R',689-700,'R',702-784 <TOK>
A;Cross-references: GB:M18633; NID:g211742; PIDN:AAA48761.1; PID:g211743
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A, Title: Repeating structure of chick tropoelastin revealed A, Reference number: A26601; MUID:87242320; PMID:3593675
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Bressan, G.M.; Argos, P.; Stanley, K.K. Biochemistry 26, 1497-1503, 1987
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                                                                                                                                                                                                       F;773-779/Disulfide bonds: #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                               A; Accession: A27264
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                                                                                                                                                                                                                                                                                                       Superfamily: elastin
                                                                                                                                                       Query Match
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25
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                                                  ω
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                                                                                                                          Similarity
     GGVPGAIPGGGVPGGGFPPGAGVGGLGAGLGAGLGAGGKPLKPGVSGLGGLGPLGLQPGA
                                  GGVPGAIP-GGVPGGVFYPGAGLGALG----GGALGPGGKPLKPVPGGLAG-----
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                                                                                                  59.2%;
ilarity 61.6%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                          Score 310; DB 2
Pred. No. 1.2e-1
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Pred. No. 1e-16;
4; Mismatches 2
                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                          from chick aorta PMID:3502711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PMID:2841924
                                                                                                                                                       DB 2;
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                                                                                                                                                  Length 784
                                                                                                     Indels
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                                                       48
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glycine-rich protein F19B15.50 - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 11-Jun-1999 #sequence_revision 11-Jun-1999 #text_change 09-Jul-2004
C;Accession: T08957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Gene: ATSP:F19B15.50
A;Map position: 4
C;Superfamily: glycine-rich cell wall structural protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R; Bevan, M.; Hilbert, H.; Braun, M.; Holzer, submitted to the Protein Sequence Database, I
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A; Accession: T08957
000
            hypothetical protein [imported] - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change
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                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: DNA
A; Residues: 1-783 < MUR>
                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Reference number: A; Accession: T35389
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A; Residues: 1-158 <BEV>
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                                                                E86179
                                                                                   RESULT 9
                                                                                                                                                                                                                                                                                   Query Match
Best Local
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Best Local :
Accession:
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                                                                                                                                   417 GGVGPGGAGSGGVGPGGAGSGGV-GPGGAG
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                                                                                                                                                                                                 PAVTFPGALVPGGVADAAAYKAAKAGAGLGGVPGVGGLG
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                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                   6
                                                                                                                                                                                                                                                                   Score 148; DB 2; 1
Pred. No. 0.00075;
6; Mismatches 39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 148; DB 2; Length 158; Pred. No. 0.0002;
                                                                                                                                                                                                                                                                                                                                                                                    EMBL: AL079348; PIDN: CAB45488.1; GSPDB: GN00070; SCOEI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -AGGLGGIGGVGGLG 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          , E.; Brandt,
May 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22;
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                                                                                                                                   455
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                                                                                                                                                                                                                                                                   Indels 10;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28;
                   09-Jul-2004
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                                                                                                                                                                                                                                                                   Gaps
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A; Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khay C.A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Mait Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H. A; Atthors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; Sker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W. A; Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis. A; Reference number: A86141; MUID:21016719; pMID:11130712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Species: hūman herpesvirus 4, Epstein-Barr virus (;Date: 25-Feb-1995 #sequence revision 25-Feb-1985 #text_chan C;Date: 25-Feb-1995 #sequence revision 25-Feb-1985 #text_chan C;Accession: C43043; S42440, Ã03773; S33021 R;Bankier, A.T.; Deininger, P.L.; Farrell, P.J.; Barrell, B.G wol. Biol. Med. 1, 21-45, 1983 A;Title: Sequence analysis of the 17,166 bp EcoRI fragment C A;Reference number: A93065; MUID:85035713; PMID:6092825 A;Accession: C43043
                                                                                                                                                                                                                                                                                                                                                                                                         A/Contents: annotation; protein coding region
R:Sample, J.; Hummel, M.; Braun, D.; Birkenbach, M.; Kieff, E.
Proc. Natl. Acad. Sci. U.S.A. 83, 5096-5109, 1986
A;Title: Nucleotide sequences of mRNAs encoding Epstein-Barr virus
A;Reference number: S424440; MUID:86259739; PMID:3460083
A;Accession: S42440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Molecule type: DNA
A;Residues: 1-641 <BANS
A;Residues: 1-641 <BANS
A;Cross-references: UNIPROT:P03211; EMBL:V01555; NID:g59074; PIDN:CAA24816.1; PID:g13348
A;Cross-references: strain B95-8
A;Experimental source: strain B95-8
R;Baer, R; Bankter, A.T.; Biggin, M.D.; Deininger, P.L.; Farrell, P.J.; Gibson, T.J.; P. Nature 310, 207-211, 1984
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Map position:
C; Superfamily:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Theologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White, O.; Alonso, Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; Dewar, K.; ansen, N.F.; Hughes, B.; Huizar, L.
Nature 408, 816-820, 2000
                                                                                                                                                                                                                             A;Cross-references: EMBL:M13941; NID:g330399; PIDN:AAA45889.1; PID:g555157
C;Superfamily: Epstein-Barr virus nuclear antigen
C;Keywords: DNA binding; transcription regulation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A,Title: DNA sequence and expression of the B95-8 Epstein-Barr virus A;Reference number: A03794; MUID:84270667; PMID:6087149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C; Genetics:
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A; Residues: 1-212 <STO>
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                                                                                                                                                                                                                                                                                                                           A; Molecule type: mRNA
A; Residues: 1-66 < SAM>
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                                                                                                                                                                                                                                                                                                                                                                                      A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nuclear antigen EBNA1 - human herpesvirus 4
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   43;
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                                              3 GGVPGAIPGGVPGGVFYPGAGLGALGGGALGPGGKPLKPVPGGLAGAGLGAGLGAFPAVT
                                                                                                                                            Similarity
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                                                                                                                Score 144.5; DB Pred. No. 0.0011; 0; Mismatches 4
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Pred. No. 0.00033;
5; Mismatches 27;
                                                                                                                                                                     DB 1;
                                                                                                                   47;
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- AGAGGAGAGGAGGAGGAG
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                                                                                                                   7;
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Maiti, R.; Marziali,
                                                                                                                   Gaps
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heat shock related protein - Plasmodium berghei
C;Species: Plasmodium berghei
C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 09-Jul-2004
C;Accession: T10455
R;Wiser, M.F.; Jennings, G.J.; Uparanukraw, P.: van Belkum A noorn t
                                                                                                                                                                                                                                                                                                                                                                                                                            A;Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G. A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complete A;Reference number: A70500; MUID:98295987; PMID:9634230 A;Accession: E70895
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A;Cross-references: GB:AL021897; GB:AL123456; NID:g3256022; A;Experimental source: strain H37Rv
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A;Experimental source: strain ANKA
C;Keywords: molecular chaperone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Status: preliminary; translated from GB/EMBL/DDBJA;Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Wiser, M.F.; Jennings, G.J.; Uparanukraw, P.; van Belkum, A.; Doorn, L.J.; Kum Mol. Biochem. Parasitol. 83, 25-33, 1996
A;Title: Further characterization of a 58 kDa Plasmodium berghei phosphoprotein A;Reference number: Z17026; MUID:97164116; PMID:9010839
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                                                                                             TFPGALVPGGVADAAAYKAAKAG-----AGLGGVPGVGGLG 98
                                                                                                                                                                   GGVPGAIPG-GVPGGVFYPGAGLGALGGGALGPGGKPLKPVPGGLAGAGLGAGLGAFPAV
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37.2%;
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39.4%;
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Pred. No. 0.0013;
                                                                                                                                                                                                                         Score 136.5; DB Pred. No. 0.0051;
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                                                                                                                                                                     61
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hypothetical glycine-rich protein Rv3508 C;Species: Mycobacterium tuberculosis
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R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Ho Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S. Nature 393, 537-544, 1998

A;Authors: Sqares, R.; Sulston, J.B.; Taylor, K.; Whitehead, S.; Barrell, B.G. A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complet A;Reference number: A70500; MUID:98295987; PMID:9634230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hypothetical glycine-rich protein Rv3514 - Mycobacterium tuberculosis (strain C;Species: Mycobacterium tuberculosis C;Date: 17-Jul-1998 #sequence_revision 17-Jul-1998 #text_change 20-Jun-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Title: Sequence and analysis of chromosome 2 of the plant Arabidopsis thaliana A;Reference number: A84420; MUID:20083487; PMID:10617197 A;Accession: G84687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Lin, X.; Kaul, S.; Rounsley, S.D.; Shea, T.P.; Benito, M.I.; Town, C.D.; Pujii, C.Y.; M.; Koo, H.; Moffat, K.S.; Cronin, L.A.; Shen, M.; VanAken, S.E.; Umayam, L.; Tallon, L. euss, D.; Nierman, W.C.; White, O.; Eisen, J.A.; Salzberg, S.L.; Fraser, C.M.; Venter,
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C;Date: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change 09-Jul-2004
C;Accession: G84687
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A;Cross-references: GB:AL022022; GB:AL123456;
A;Experimental source: strain H37Rv
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                                                                                                                                                                                                                                                                                   ;Superfamily:
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Mycobacterium tuberculosis

(strain

H37RV)

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C;Accession; F70806, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Gordon, S. T.; Broach, R.; Davies, R.; Seeger, K.; Skelton, S.; Squares, S.; Squares, R.; Sulston, J.E.; Teylor, K.; Whitehead, S.; Barrell, B.G. A.; Title: Deciphering the biology of Mycobacterium tuberculosis from the complete genome A; Residues preliminary nucleic acid sequence not shown; translation not shown A; Status; preliminary nucleic acid sequence not shown; translation not shown A; Residues; 1191 (COL) (CO
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Copyright (c) 1993 - 2005 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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"Structure of the 3' region of the human elastin gene: of Alu repetitive sequences and few coding sequences.";
Connect. Tissue Res. 16:197-211(1987).
EMBL; M17282; AAC98393.1;
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Indik Z., Yeh H., Ornstein-Goldstein N., Sheppard P.,
Rosenbloom J.C., Petichnen L., Rosenbloom J.;
Rosenbloom J.C., Petichnen L., Rosenbloom J.;
"Alternative splicing of human elastin mRNA indicated
"Alternative splicing of human elastin mRNA indicated
analysis of cloned genomic and complementary DNA.";
Proc. Natl. Acad. Sci. U.S.A. 84:5680-5684(1987).
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Q7Z3F5;
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                Fazio M.J., Olsen D.R., Kauh E.A., Baldwin C.T., Indik Z., Ornstein-Goldstein N., Yeh H., Rosenbloom J., Uitto J.; "Cloning of full-length elastin cDNAs from a human skin fibroblast recombinant cDNA library: further elucidation of alternative spliciutilizing exon-specific oligonucleotides."; J. Invest. Dermatol. 91:458-464(1988).
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                                                                                                                                                                                                                                             Indik Z., Yeh H., Ornstein-Goldstein N., Sheppard P., Rosenbloom J.C., Peltonen L., Rosenbloom M.G., Peltonen E., Rosenbloom M., William of the Manalysis of cloned genomic and complementary DNA, "; proc. Natl. Acad. Sci. U.S.A. 84:5680-5684 (1987).
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Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9606;
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Name=DKFZp686F06102;
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                                                                                                                                                        TISSUE=Skin fibroblast;
MEDLINE=89009960; PubMed=3171221;
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TISSUE=Human fetal
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"Isolated supravalvular aortic stenosis: functional haploinsufficiency
of the elastin gene as a result of nonsense-mediated decay.";
Hum. Genet. 106:577-588(2000).
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J. Biol. Chem. 274
[6]
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                                                                                                                          -I- PTM: The crosslinks are made of deaminated Lys.
-I- DISEASE: Defects in ELN are a cause of autosomal dominant cutis laxa [MIM:123700]. Cutis laxa is a rare connective tissue disorder characterized by loose, hyperextensible skin with decreased resilience and elasticity leading to a premature aged appearance. The skin changes are often accompanied by extracutaneous manifestations, including pulmonary emphysema, bladder diverticula, pulmonary artery stenosis and pyloric stenosis.
-I- DISEASE: Haploinsufficiency of ELN may be the cause of certain cardiovascular and musculo-skeletal abnormalities observed in Williams-Beuren syndrome (WBS) [MIM:194050]. WBS is a rare developmental disorder and a contiguous gene deletion syndrome developmental disorder and a contiguous gene deletion syndrome
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Cell 86:59-69(1996).
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MEDLINE=96291399; PubMed=8689688; DOI=10.1016/S0092-8674(00)80077-X;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TISSUE=Placenta;
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involving genes from chromosome band 7q11.23. DISEASE: Defects in ELN are the cause of supravalvular aortic stenosis (SVAS) [MMM:185500]. SVAS is a congenital narrowing the ascending aorta which can occur sporadically, as an autose dominant condition, or as one component of Williams-Beuren
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FUNCTION: Major structural protein of tissues such as aorta and nuchal ligament, which must expand rapidly and recover complete SUBUNIT: The polymeric elastin chains are cross-linked together into an extensible 3D network.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SUBCELLULAR LOCATION: Extracellular matrix of elastic fibers. ALTERNATIVE PRODUCTS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Name=1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Event=Alternative splicing; Named isoforms=2; Comment=Additional isoforms seem to exist;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IsoId=P15502-2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IsoId=P15502-1; Sequence=Displayed;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence=VSP_004243;
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RESULT Q14234 ID Q14234 ID G1 AC Q1 AC Q1 DT 01 DT 01 DT 01 DE E1 DE E1 GN Na OCC EU OCC Ma
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MIM; 123700;
MIM; 194050;
MIM; 194550;
GO; GO:000551;
GO; GO:000520;
GO; GO:000828;
GO; GO:000888;
GO; GO:0009888;
GO; GO:0009888;
GO; GO:0009888;
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Q14234;
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                   Homo sapiens (Human
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
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01-MAR-2004
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                                                                                                      Name=ELN;
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C:extracellular space; TAS.
F:extracellular matrix structural constituent;
P:cell proliferation; TAS.
P:circulation; TAS.
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Q75MU5;
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MEDLINE-87289668; PubMed=3039501;

MINDINE-87289668; PubMed=3039501;

MINDINE-87289668; PubMed=3039501;

Rosenbloom J.C., Peltonen L., Rosenbloom J.;

Rosenbloom J.C., Peltonen L., Rosenbloom J.;

"Alternative splicing of human elastin mRNA indicated analysis of cloned genomic and complementary DNA.";

Proc. Natl. Acad. Sci. U.S.A. 84:5680-5684(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      rosenation J., ornstein-Goldstein N.;
"Structure of the 3' region of the human elastin gene
of Alu repetitive sequences and few coding sequences.
Connect. Tissue Res. 16:197-211(1987).
                   MEDLINE=22737999; Pul
Hillier L.W., Fulton
                                                                                                                                      Eukaryota; Metazoa;
Mammalia; Eutheria;
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MEDLINE-87274906; PubMed-3038460;
MORTOW S.D., Cicila G.,
                                                                  SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                      Hypothetical protein
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PROSITE; PS00101; HEXAPEP_TRAN
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InterPro; IPR003979; tropoelastin.
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HI; M17265; AAC98395.1; JOINED

LI; M17267; AAC98395.1; JOINED

LI; M17268; AAC98395.1; JOINED

LI; M17268; AAC98395.1; JOINED

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GO:0005578; C:extracellular matrix (s
GO:0005578; F:extracellular matrix co
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GO:0030023;
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Ornstein-Goldstein N.;
PubMed=12853948; DOI=10.1038/nature01782;
ton R.S., Fulton L.A., Graves T.A., Pepin
C., Layman D., Maas J., Jaeger S., Walker
                                                                                                                                         Chordata;
Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              66136 MW;
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                                                                                                                                                                                                                                      ELN.
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                                                                                                                                                                                                                                                         Created)
Last sequence update)
Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 5
                                                                                                                                      Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23B7FE5B8AF85CA8 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                                                                                                                                                                                                                                         757
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RA Wylie K., Sekhon M., Becker M.C., O'Laughlin M.D., Schaller M.E.,
RA Sun H., Edwards J., Bradshaw-Cordum H., Ali J., Andrews S., Isak A.,
RA Vanbrunt A., Nguyen C., Du F., Lamar B., Courtney L., Kalicki J.,
RA Ozersky P., Bielicki L., Scott K., Holmes A., Harkins R., Harris A.,
RA Strong C.M., Hou S., Tomlinson C., Dauphin-Kohlberg S.,
RA Kozlowicz-Reilly A., Leonard S., Rohlfing T., Rock S.M.,
RA Tin-Wollam A.M., Abbott A., Minx P., Mauphin R., Strowmatt C.,
RA Tin-Wollam A.M., Abbott A., Minx P., Maupin R., Strowmatt C.,
RA Hickeille P., Miller N., Johnson D., Murray J., Woessner J.P.,
RA Hickenbotham M.T., Eldred J., Williams D., Bedell J.A., Mardis E.R.,
RA Gillett W., Zhou Y., James R., Phelps K., Iadanoto S., Bubb K.,
RA Gillett W., Zhou Y., James R., Phelps K., Iadanoto S., Bubb K.,
RA Simms E., Levy R., Clendenning J., Kaul R., Kent W.J., Furey T.S.,
RA Bailey J.A., Portnoy M.E., Torrents D., Chinwalla A.T., Gish W.R.,
RA Bailey J.A., Portnoy M.E., Torrents D., Chinwalla A.T., Gish W.R.,
RA Eddy S.R., McPhetson J.D., Olson M.V., Eichler E.E., Green E.D.,
RA Waterston R.H., Wilson R.K.;
UN [2]
RESULT 7
O1533
AC 0153
AC 0153
DT 01-3
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DT 01-8
COS HOMMO
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Best Local
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015337;
01-JAN-1998
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Submitted
                          SEQUENCE FROM N.A.

MEDLINE=97358574; PubMed=9215670; DOI=10.1093/
Li D.Y., Toland A.E., Boak B.B., Atkinson D.L.

Morris C.A., Keating M.T.;

"Elastin point mutations cause an obstructive supravalvular aortic stenosis.";

Hum. Mol. Genet. 6:1021-1028(1997).
                                                                                                                                                                                                                                                                                                                                         Elastin (Fragment)
Homo sapiens (Huma
                                                                                                                                                                                                                                                                                                                                                                                           01-JAN-1998 (TrEMBLrel.
01-JAN-1998 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted (JAN-2004) to the EMBL/GenBank/DDBJ EMBL; AC005056; AAS07435.1; -. GO: GO:0005578; C:extracellular matrix (sensu GO: GO:0005201; F:extracellular matrix structu InterPro; IPR001451; Hexapep_transf. InterPro; IPR003979; tropoelastin.
                                                                                                                                                                                                                                               NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hypothetical
SEQUENCE 75
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PROSITE; PS00101; HEXAPEP_TRANSFERASES; UNKNOWN_1.
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  Mol. Genet.
U93037; AAI
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                                                                                                                                                                                                                                                                                  Chordata;
Primates;
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Pred. No.
                                                                                                                                                                                                                                                                                     Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                              sequence update)
annotation update)
                                                                                                                                         DOI=10.1093/hmg/6.7.1021;
Atkinson D.L., Ensing G.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               602
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (sensu Metazoa);
structural const:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6e-25;
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RESULT
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REMBL; U93035; AAB65620.1; JOINED.

REMBL; U93036; AAB65620.1; JOINED.

RGO; GO:0005578; C:extracellular matrix (sensu Metazoa); IEA.

RGO; GO:0005201; F:extracellular matrix structural constituent; IEA.

RINEETPO; IPR001451; HEXAPEP_transf.

INTERPO; PR01500; TROPOELLSTIN.

RPRNITS; PR01500; TROPOELLSTIN.

PROSITE; PS01011; HEXAPEP_TRANSFERASES; UNKNOWN_1.

TRON_TER 602 602

SEQUENCE 602 AA; 51807 MW; 53B5B9A71EF04807 CRC64;
                                                                                                                                                                                                    Query Match
Best Local S
Matches 96
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; U93037; AAB65621.1; -.
EMBL; U93034; AAB65621.1; JOINED.
EMBL; U93035; AAB65621.1; JOINED.
EMBL; U93036; AAB65621.1; JOINED.
GO; GO:0005578; C:extracellular matrix
GO; GO:0005201; F:extracellular matrix;
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NON TER 1 1

NON TER 635 635

SEQÜENCE 635 AA; 55279 MW; 72950C364127B2A4 CR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=97358574; PubMed=9
Li D.Y., Toland A.E., Boak
Morris C.A., Keating M.T.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                    InterPro; IPR001451; Hexapep_transf. InterPro; IPR003979; tropoelastin. PRINTS; PR01500; TROPOELASTIN.
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Homo sapiens (Human).
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                                                                                              GALVPGGVADAAAAYKAAKAGAGLGGVPGVGGLGVS
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.E., Boak B.B., 1
                                                                                                                                                                                                  96.0%; Score 503; DI
100.0%; Pred. No. 3e:
tive 0; Mismatches
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Pred. No. 2.9e-24;
                                                                                                                                                                                                                                                                                                        72950C364127B2A4 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DOI=10.1093/hmg/6.7.1021;
Atkinson D.L., Ensing G.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                structural constituent;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (sensu Metazoa);
                                                                                                                                                                                                                         3e-24;
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06ZUN2;
05-JUL-2004
05-JUL-2004
05-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Q7Z316 PRELIMINARY; PKT; GO, A., Q7Z316; Q7Z316; Created) 01-OCT-2003 (TrEMBLrel. 25, Last sequence update) 01-OCT-2003 (TrEMBLrel. 25, Last annotation update) 01-MAR-2004 (TrEMBLrel. 26, Last annotation update).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oshima A., Takahashi-Fujii A., Tanase T., Imose N., Takeuchi K., Arita M., Musashino K., Yuuki H., Hara H., Sugiyama T., Irie R., Otsuki T., Sato H., Wakamatsu A., Ishii S., Yamamoto J., Isono Y., Kawai-Hio Y., Saito K., Nishikawa T., Kimura K., Yamashita H., Kawai-Hio Y., Saito K., Nishikawa T., Kimura K., Yamashita H., Watsuo K., Nakamura Y., Sekine M., Kikuchi H., Kanda K., Wagatsuma Murakawa K., Kanehori K., Sugiyama A., Kawakami B., Suzuki Y., Sugiyama A., Kawakami B., Suzuki Y., Sugiyamo S., Nagahari K., Masuho Y., Nagai K., Isogai T.; Submitted (JUL-2003) to the EMBL/GenBank/DDBJ databases.
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein FLJ43523.
Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; AK125511; BAC86188.1; -. GO; GO;0005578; C:extracellular matrix (sensu Metazoa); GO; GO:0005201; F:extracellular matrix structural constituterbro; IPR003979; tropoelastin.
PRINTS; PR01500; TROPOELASTIN.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE-Placenta;
Oshima A., Takahashi-Fujii A.,
                                                                                                                                                                            Submitted (JUN-2003) to the EMBL/GenBank/DDBJ EMBL; BX538199; CAD98065.1; -. InterPro; IPR001451; Hexapep transf. PR001TE; PS00101.1** EXAPEP_TRĀNSFERASES; UNKNOW HYDOTHETICAL PROSTERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hypothetical protein DKFZp686021208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
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                                                                                                                                    Hypothetical protein.
NON_TER 1
                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                     Han M.,
                                                                                                                                                                                                                                                                                                                                                           rissum=Human rectum tumor;
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                                                                                                                                                                                                                                                                                                  Han M., Wiemann S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      9
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                         Similarity
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Conservative
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                                                                                                                                                                                                                                                                                                                              Mewes H.W.,
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No. 7.8e-21;
                            Pred.
                                                   Score 448;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Craniata; Vertebrata; I Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRT;
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constituent;
                                                                                                                                                                                                                                                                                                                                 Osanger
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AC P5432
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Matches 86
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Nagahari K., Sugano S., Isogai T.;

Submitted (MAR-2002) to the EMBL/GenBank/DDBJ databases.

EMBL; AKO75554; BAC11696.1; -.

HSSP; P50099; 1ZFJ.

GO; GO:0005278; C:extracellular matrix (sensu Metazoa); I

GO; GO:0005201; F:extracellular matrix structural constit

InterPro; IPR001451; Hexapep_transf.

InterPro; IPR003979; tropoelastin.

PRINTS; PR01500; TROPOELASTIN.

PROSITE; PS00101; HEXAPEP TRANSFERASES; UNKNOWN 1.

PROSITE; PS00101; HEXAPEP TRANSFERASES; UNKNOWN 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        P54320;
01-OCT-1996
01-OCT-1996
25-OCT-2004
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                                                                                             STRAIN=BALB/c; TISSUE=Lung;
MEDLINE=95130069; PubMed=7829060;
Wydner K.S., Sechler J.L., Boyd C
"Use of an intron polymorphism to
mouse chromosome 5 in a region of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-OCT-2002 (TrEMBLrel. 22, Creat
01-OCT-2002 (TrEMBLrel. 22, Last
01-MAR-2004 (TrEMBLrel. 26, Last
                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-OCT-1996 (Rel. 34, Created)
01-OCT-1996 (Rel. 34, Last sequence update)
25-OCT-2004 (Rel. 45, Last annotation update)
Elastin precursor (Tropoelastin).
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                                  chromosome 7.";
Genomics 23:125-131(1994)
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                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=10090;
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   Major structural
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                                                                                                                                                                                                                                                                                                                                                                     ; Chordata;
; Rodentia;
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, Wakamatsu A., Nagai
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Last annotation update)
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linkage conservation with
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No. 3.7e-20;
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01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-MAR-2003 (TrEMBLrel. 28, Last sequence update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
Mus musculus 3 days neonate thymus cDNA, RIKEN full-length enriched
library, clone:A630042I19 product:elastin, full insert sequence
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                      SEQUENCE FROM N.A.
STRAIN-C57BL/6J; TISSUE-Thymus;
The FANTOM Consortium,
the RIKEN Genome Exploration Re
                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
STRAIN=C57BL/6J; TISSUE=Thymus;
MEDLINE=21085660; PubMed=11217851;
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STRAIN=C57BL/6J; TISSUE=Thymus;
MEDLINE=99279253; PubMed=10349636;
                                                                                                                                                                               RIKEN FANTOM Consortium; "Functional annotation of a Nature 409:685-690(2001).
                                                                                                                                                                                                                                                                                                                                                                                                   Carninci P., Hayashizaki Y.; "High-efficiency full-length cDNA Meth. Enzymol. 303:19-44(1999).
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SIGNAL 1 27 Potent
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SUBCELLULAR LOCATION: Extracellular matrix of elastic f PTM: The crosslinks are made of deaminated Lys.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SWISS-PROT entry is copyright. It is produced through a collaboration
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   mouse
                               Exploration Research
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Pred. No. 3.1e-16;
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Sciurognathi;
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h Group
based o
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thi; Muridae; Murinae; Mus
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p Phase
on func
hase I & II
functional
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SEQUENCE Nature 60,770

420:563-573 full-length cDNAs.";

(2002).

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SEQUENCE FROM N.A.

STRAIN=FVB/N; TISSUE=Mammary tumor. C3;

STRAIN=FVB/N; TISSUE=Mammary tumor. C3;

XX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

XX MILSCHUN C., Feingold E.A., Grouse L.H., Schaefer C.F., Bhat N.K.,

XX Allschul S.F., Zeeberg B., Buertow K.H., Schaefer C.F., Bhat N.K.,

XX Allschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

XX Allschul R.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,

XX Allschul R.F., Jordan M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

XX Allschul R.F., Jordan N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

XX Allschul R.F., Moorley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

XX Allschul S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

XX Allslon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,

XX Allschul R.F., Foung A.C., Shevchenko Y., Bouffard G.G.,

XX Millalon D.K., Muzny D.M., Green E.D., Dickson M.C.,

XX Allschul R.F., Tourdan J.W., Green E.D., Dickson M.C.,

XX Allschul R.F., Schalska U., Smailus D.E., Schnerch A., Schein J.E.,

XX Allschul R.F., Schalska U., Smailus D.E., Schnerch A., Schein J.E.,

XX Allschul R.F., Schalska U., Smailus D.E., Schnerch A., Schein J.E.,
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(STRAIN-C578L/6J; TISSUE-Thymus;

(STRAIN-C578L/6J; TISSUE-Thymus;

(MEDIINE-20530913; PubMed=11076861; DOI=10.1101/gr.152600;

(MEDIINE-20530913; PubMed=11076861; DOI=10.1101/gr.152600;

(MEDIINE-20530913; PubMed=11076861; DOI=10.1101/gr.152600;

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(MEDIINE-20530913; PubMed=10.76861; DOI=10.1101/gr.152600;

(MEDIINE-20530913; PubMed=110.1101/gr.152600;

(MEDIINE-20530913; PubMed=10.1101/gr.152600;

(MEDIINE-20530913; PubMed=10.1101/gr.152600;

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Genome F
         MGD; NGC
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Konno H., Okazaki Y., Muramatsu M., Hayashiaki Y.;
"Normalization and subtraction of cap-trapper-selected
prepare full-length cDNA libraries for rapid discovery
                                     Submitted (APR-2003) to the EMBL; AK041860; BAC31084.1; EMBL; BC051649; AAH51649.1; MGD; MGI:95317; Eln.
                                                                                                                                                                                                                                                                                                                     Jones S.J., Marra M.A.; "Generation and initial analysis of more than 15,000
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STRAIN=C57BL/6J;
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                                                                                                                                                    Strausberg R.;
                                                                                                                                                                             SEQUENCE FROM N.A.
STRAIN=FVB/N; TISSUE=Mammary
                                                                                                                                                                                                                                                                         mouse cDNA sequences.";
c. Natl. Acad. Sci. U.S.A.
         GO:0007519;
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Res. 10:1617-1630(2000).
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6J; TISSUE=Thymus;
AAH51649.1; -.
Eln.
P:myogenesis; IMP
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Matches 77
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PRINTS; PR01500; TROPOELASTIN.
                                                     Q6ZWJ6;
05-JUL-2004
05-JUL-2004
05-JUL-2004
                                05-JUL-2004 (TrEMBLrel. 27, Created)
05-JUL-2004 (TrEMBLrel. 27, Last seq
05-JUL-2004 (TrEMBLrel. 27, Last ann
Hypothetical protein FLJ16246.
Homo sapiens (Human)
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01-OCT-2002 (TrEMBLrel. 22, Last sequence up
01-MAR-2004 (TrEMBLrel. 26, Last annotation
Hypothetical protein PSEC0191.
Homo sapiens (Human).
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         Homo sapiens (Human
Eukaryota; Metazoa;
Mammalia; Eutheria;
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NON_TER 472 472
SEQUENCE 472 AA; 42265 MW; 8D7A0F3A
                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE=Whole embryo;
Ota T., Nishikawa T., Suzuki Y.,
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01-OCT-2002
                                                                                                    9 PMZ 9 D
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                                                                                                                                                              IPGGVVGAGPAAAAAAAAKAAKYGAAAGLVPGGPGFG
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72.6%;
                     Chordata;
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79.4%;
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Pred. No. 2.
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1; Mismatches
          Craniata; V
Catarrhini;
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                       Vertebrata; Euteleostomi;
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Nakamura Y.
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Primates;

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A Tanigami A., Fujiwara T., Shibahara T., Goto Y., Hirao M., Shimizu F.,

A Tanigami A., Fujiwara T., Shibahara T., Ozaki K., Sugiyama T.,

A Irie R., Otsuki T., Sato H., Wakamatsu A., Ishii S., Yamamoto J.,

A Irie R., Otsuki T., Sato H., Wakamatsu A., Ishii S., Yamamoto J.,

A Irie R., Otsuki T., Sato K., Nishikawa T., Kimura K.,

Yamashita H., Matsuo K., Nakamura Y., Sekine M., Kikuchi H., Kanda K.,

Yamashita H., Matsuo K., Nakamura Y., Sekine M., Kikuchi H., Kanda K.,

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Yamashita H., Matsuo K., Nakamura Y., Sekine M., Kikuchi H., Kanda K.,

Yamashita H., Matsuo K., Nakamura Y., Sekine M., Kikuchi H., Kanda K.,

Yamashita H., Yamashita H., Suzuki Y., Sugano S., Nagahari K.,

Yamashita H., Yawashita H., Suzuki Y., Sugano S., Nagahari K.,

Yamashita H., Yawashita H., Wazuki Y., Sugano S., Nagahari K.,

Yamashita H., Yawashita H., Wazuki Y., Sugano S., Nagahari K.,

Yamashita H., Yawashita H., Wazuki Y., Sugano S., Nagahari K.,

Yamashita H., Yawashita H., Wazuki Y., Sugano S., Nagahari K.,

Yamashita H., Yawashita H., Wazuki Y., Sugano S., Nagahari K.,

Yamashita H., Yawashita H., Wazuki Y., Sugano S., Nagahari K.,

Yamashita H., Yawashita H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 67.0
Best Local Similarity 57.4
Matches 81; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    InterPro; IPR001451; Hexapep_transf.
InterPro; IPR003979; troppelastin.
PRINTS; PR01500; TROPOELASTIN.
PROSITE; PS00101; HEXAPEP TRANSFERASES; UNKNOWN_1.
SEQUENCE 570 AA; 48941 MW; 3117B028D06D4F7B CRC64;
                                                                                                                                                                                      145 AAAKFGAGAAGVLPGVGGAGV 165
                                                                                                                                                                                                                                        80 KAAKAGAGLGGV-PGVGGLGV 99
                                                                                                                                                                                                                                                                                                                                                          87 -- PGVGPFGGPQPGVPLGYPIKAPKLPGYGPGGVAGAAGKAGYPTGTGVGPQAAAAAAK 144
                                                                                                                                                                                                                                                                                                                                                                                                                              62 TFPG------ALVPGGVAD------AAAAY 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27 GGVPGAIPGGVPGGVFYPGAGLGALGGGALGPGGKPLKPVPGGLAGAGLGAGVGGAFAGI 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3 GGVPGAIPGGVPGGVFYPGAGI.GAILGGGAILGPGGKPLKPVPGGLAGAGI.GAGI.GAFPAV 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      67.0%; Score 351; DB 2; Length 570; 57.4%; Pred. No. 9.9e-15; tive 2; Mismatches 12; Indels 46; Gaps
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Database
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Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                            hits satisfying chosen parameters:
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501
                                                                                                                                                                                                                                                                                                                                                                                                                                                August 10, 2005, 07:16:57 ; Search time 120.095 Seconds (without alignments) 325.267 Million cell updates/sec
                                                                                                                                                                                                                                                                                                          2105692 seqs, 386760381 residues
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Gapop 10.0 ,
                                                                                                             A_Geneseq_16Dec04:*
1: geneseqp1980s:*
                                                                                                                                                              Maximum Match 100%
Listing first 45 s
                                                                                                                                                                                                                                                                                                                                                                                          VAAAAKSAAKVAAKAQLRAA.....QHLPSTPSSPRVPGALAAAK 101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GenCore version
Copyright (c) 1993 - 2005
                geneseqn1990s:*
geneseqp2000s:*
geneseqp2001s:*
geneseqp2002s:*
geneseqp2003as:*
geneseqp2003bs:*
geneseqp2004s:*
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Compugen Ltd.
                                                                                                                                                                                                                                                                             2105692
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

### SUMMARIES

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Result No.	Score	Query Match	Length	BB	ID	Description
۲	501	100.0	730	N	AAW46315	Human e
N	501		731	ω	AAY69068	Amino
w	501	100.0	731	4	AAB66657	Human
4.	501	100.0	731	σ	ABU08725	Abu08725 Human ela
ۍ	501	•	731	7	ADL96420	Ad196420 Human ela
o	501	•	733	N	AAR56653	Aar56653 Synthetic
7	501	•	733	N	AAY01301	Aay01301 Amino aci
80	501	•	757	7	ABG75223	Abg75223 Human tro
9	501	100.0	757	7	ADP65160	Human
10	424	84.6	216	N	AAY01310	Aay01310 Human tro
11	308	61.5	200	N	AAY01305	Aay01305 Human tro
12	304.5	60.8	617	7	ADB64761	Adb64761 Human pro
13	304.5		660	N	AAY01303	Aay01303 Human tro
14	304.5	•	663	7	ADM03792	Adm03792 Human pro
15	304.5	60.8	692	7	ADE40134	Human
16	304.5		698	ນ	AAY01302	Human
17	304.5		698	w	AAY69069	Aay69069 Amino aci
18	304.5	•	712	w	AAB08630	Amino
19	304.5	•	730	w	AAB08631	Aab08631 Fusion pr
20	304.5	60.8	730	ហ	AAO17360	Aao17360 Human ela
21	304.5	•	730	œ	ADQ19747	Adq19747 Human sof
22	304.5	60.8	870	7	ADE08527	Ade08527 Novel pro
23	303.5		570	ထ	ADQ65871	
24	301.5	60.2	711	J	ADE40132	Ade40132 Human NOV
25		,		,		

This represents the human elastin sequence containing the minimal functional unit (MFU)-1 polypeptide of the invention. This MFU-1 is a polypeptide that has at least 3 beta-sheet/ beta-turn structures, but is not a naturally occurring fibrous protein. Each beta-sheet structure has 3-7 (preferably 5-7) amino acids and the MFU polypeptide may include at least 1 amino acid that can take part in crosslinking. The polypeptide can also be derived from the sequences of animal elastin, lamprin and

New non-natural polypeptide with multiple beta-sheet, beta-turn structures - particularly based on human elastin, useful for coating prostheses, as wound dressings, etc., allows ingrowth of cells.

Claim 5; Fig 1B; 39pp; English.

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	0
154	154	154	154	154	154.5	155	155	158	158	159	159	159	170	174	191	227.5	227.5	249.5	100
30.7	30.7	30.7	30.7	30.7	30.8	30.9	30.9	31.5	31.5	31.7	31.7	31.7	33.9	34.7	38.1	45.4	45.4	49.8	((, )
884	884	246	246	244	235	877	877	287	223	281	217	34	34	34	171	183	183	864	U / F
N	N	œ	œ	œ	N	N	N	N	N	N	N	Ц	N	N	w	w	N	7	•
AAW09213	AAR80341	ADR70474	ADR70476	ADR70475	AAR80349	AAW49724	AAR80335	AAW49737	AAR80348	AAW49736	AAR80347	AAP71455	AAY01309	AAY01308	AAY69137	AAY69138	AAY01311	ADE56670	
Aaw09213	Aar80341	Adr70474	Adr70476	Adr70475	Aar80349	Aaw49724	Aar80335	. Aaw49737	Aar80348	Aaw49736	Aar80347	Aap71455	Aay01309	Aay01308	Aay69137	Aay69138	Aay01311	Ade56670	The Paris of the P
SELP8K po	Protein p	Silk-elas	Silk-elas	Silk-elas	Protein p	Protein p	Protein p	Protein p	Protein p	Protein p	Protein p	Sequence	Human tro	Human tro	Amino aci	Amino aci	Human tro	Rat Prote	

# ALIGNMENTS

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RESULT 1
AAW46315
Key
Protein
                                                                                                                                                                                                                                                                                          23-JUL-1998
                                                                                                                                                                                                                                                                                                                     AAW46315 standard; protein;
                                                                                                                                              07-AUG-1996;
07-AUG-1997;
                                                                                                                                                                    07-AUG-1997;
                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                       beta-sheet; coating;
                                                                                                                                                                                                                                                              MFU-1; minimal functional unit; elastin; human; fibrous protein;
                                                                                                                                                                                                                                                                            Human elastin
                                                                                                                                                                                                                                                                                                        AAW46315;
                                                                                               WPI; 1998-145551/13.
                                                                                                            Rothstein A,
                                                                                                                         (PROT-) PROTEIN SPECIALTIES LTD. (HOSP-) HOSPITAL FOR SICK CHILDREN.
                                                                                                                                                                                  12-FEB-1998.
                                                                                                                                                                                                WO9805685-A2
                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                            Keeley FW,
                                                                                                                                                                                                                                                                           containing non-natural polypeptide MFU-1 sequence
                                                                                                                                               96US-0023552P.
97US-00911364.
                                                                                                                                                                    97WO-CA000560
                                                                                                                                                                                                                    Location/Qualifiers 374. .499
                                                                                                                                                                                                              /note=
                                                                                                                                                                                                                                                       wound dressing.
                                                                                                                                                                                                              "MFU-1 polypeptide"
                                                                                                             Rothstein SJ;
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RESULT 2
AAY69068
ID AAY6
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           spider silk protein. The MFU polypeptides are self-aligning peptides having the same primary structure as part of a natural fibrous protein. They are used to coat prostheses made of animal or synthetic material or metal, particularly for use as blood vessel or heart valve replacements, wound or burn dreesings, or stents. They can be used in cosmetic, elastic or high-tensile strength materials, e.g. ropes or parachnte cord. Prostheses based on the MFU allow penetration of endothelial cells, so become permanent, living, tissue replacements. The MFU polypeptides have better biocompatibility than known elastin-based materials. They are well defined, homogeneous material and are easier to manipulate and produce than full-length elastins. They are non-thrombogenic and non-immunogenic. Materials can be made from 2 or more different MFU polypeptides to allow properties to be tailored for particular applications, e.g. combining the high extensibility of elastin and the high tensile strength of spider
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tropoelastin; derivative; proteolysis; protease; antiwrinkle; hand lotion; bulking agent; chemotaxis; proliferation; growth inhibition; peptidomimetic; lung damage; elastin; cancer; metastasis; blood clotting.
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Cleavage-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-MAY-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY69068 standard; protein; 731 AA
cell growth.
                            New derivatives of tropoelastin useful, e.g.
                                                                                      N-PSDB; AAZ61146.
                                                                                                                WPI; 2000-182399/16
                                                                                                                                                                     Weiss AS
                                                                                                                                                                                                                                                                          17-JUL-1998;
                                                                                                                                                                                                                                                                                                                               19-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                    27-JAN-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200004043-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cleavage-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          leavage-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        730 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence of a human tropoelastin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
441..442
503..504
515..516
564..565
                                                                                                                                                                                                                                                                          98AU-00004723
                                                                                                                                                                                                                                                                                                                               99WO-AU000580
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 501; DB 2;
Pred. No. 4.4e-34;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           splice
                              for inducing chemotaxis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 730;
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Disclosure; Page 107-109; 136pp; English

The present invention relates to a minimal functional unit (MFU) of human elastin polypeptide. This protein is useful in a cosmetic material or a prosthetic material such as prosthesis for blood vessel replacements, for

1; Fig

1; 39pp; English.

Novel polypeptides that comprise three beta-sheet/beta-turn structures and are not naturally occurring fibrous protein, used to produce prosthesis suitable for implantation into humans, and cosmetic materials.

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RESULT 3
AAB66657
ID AAB6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC The present sequence represents a human tropoelastin splice form. The specification describes tropoelastin derivatives, in which a subsequence CC has been mutated so that susceptibility to proteolysis is reduced or CC eliminated, or a subsequence has been inserted so that susceptibility to proteolysis is increased. The derivatives have with reduced CC susceptibility, and can be used where the wild-type protein would be CC degraded too easily, e.g. in contact with serum or wound exudate. The CC tropoelastin derivatives provide competitive inhibition of protease CC activity. The tropoelastin derivatives, and other polypeptides containing CC in human or veterinary medicine, cosmetics (e.g. antiwrinkle or hand CC useful for proliferation or growth inhibition, particularly of smooth CC muscle cells, epithelial or endothelial cells, fibroblasts, osteocytes, CC chondrocytes and platelets. Peptidominetics that mimic the protease CC cleavage site in tropoelastin derivatives are competitive inhibitors of CC useful, for inhibiting or controlling localized growth of cancers or CC elastin, for inhibiting or controlling localized growth of cancers or CC metastases, or to limit protease activity that causes blood clotting or metastases, or to limit protease activity that causes blood clotting or controlling localized growth of cancers or CC metastases, or to limit protease activity that causes blood clotting tropoelasting causes blood clotting or controlling localized growth of cancers or CC metastases, or to limit protease activity that causes blood clotting tropoelasting causes blood clotting controlling localized growth of cancers or contro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 101; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 731
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAB66657;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB66657 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                        29-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                   29-JUN-2000; 2000WO-US017829
                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-JAN-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Minimal function unit; MFU; human; elastin prosthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human elastin protein without signal peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-APR-2001
                                                                                                                                                                                                                      WPI; 2001-102886/11.
                                                                                                                                                                                                                                                              Rothstein A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200100666-A2
                                                                                                                                                                                                                                                                                                         (HSCR-)
                                                                                                                                                                                                                                                                                                                                (PROT-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     498 VAAAAKSAAKVAAKAQLRAAAGLGAGIPGLGVGVGVPGLGVGAGVPGLGVGAGVPGFGAG
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                                                                                                                                                                                                                                                                                                       PROTEIN SPECIALTIES LTD HSC RES & DEV LP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; !
iilarity 100.0%; !
Conservative 0;
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Pred. No. 4.4e-34;
0; Mismatches 0;
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RESULT 4
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           The invention describes a polypeptide (I) comprising a minimal functioning unit (MFU) which is present in the sequence of human elastin and comprised of at least three beta-sheet/beta-turn structures, and at least one amino acid residue that participates in cross-linking, and not a naturally occurring fibrous protein. The MFU material can be used to construct human elastin-like prostheses such as tubes for blood vessel replacement, heart replacement valves and sheets for other uses such as burns or wounds to promote healing. MFUs can be co-aggregated with other proteins, for e.g. collagen, to provide prosthesis material that resembles the natural structural materials of the body. MFUs modeled on lamprin and other fibrous proteins e.g. spider silk, can be used to make a variety of materials, for a number of different applications, for e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; elastin; minimal functioning unit; MFU; beta-sheet/beta-turn structure; fibrous protein; prosthesis; blood vessel replacement; heart replacement valve; burn; wound; lamprin; spider silk; cord; rope; parachute; cosmetic; platelet binding inhibitor; platelet activation inhibitor; non-thrombogenic; cell infiltration; non-immunogenic; biocompatible; high tensile strength; elasticity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     heart valve replacement, tissue replacement, for covering burns, covering wounds and stents
                                                                                                                                                                                                                                              Novel polypeptide useful in prosthesis, has a secondary characterized by three beta-sheet/beta-turn structures, naturally occurring fibrous protein.
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07-AUG-1997;
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                                                                                                                                                                                                                  Claim 1; Fig 1B; 21pp; English.
                                                                                                                                                                                                                                                                                                               WPI; 2003-391056/37
                                                                                                                                                                                                                                                                                                                                          Rothstein A,
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and ropes
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97US-00911364.
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use in parachutes and in
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Best Local S
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07-AUG-1997;
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blood vessel; wound; burn healing; collagen.
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  WPI; 2003-898105/82
                                                    Rothstein A,
                                                                                                      (ROTH/) ROTHSTEIN A. (KEEL/) KEELEY F. (ROTH/) ROTHSTEIN S.
                                                                                                                                                                                                                                                                                                                       28-SEP-2001; 2001US-00964662
                                                                                                                                                                                                                                                                                                                                                                                                                                US2003166846-A1.
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97US-00911364.
99US-00340736.
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Pred. No. 4.4e-34;
; Mismatches 0;
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RESULT 6
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention describes a polypeptide that comprises three beta sheet/beta turn structures and that is not a naturally occurring fibrous protein. The invention also describes a prosthesis comprising an animal, metal or synthetic material, where the surface is coated with the
                                                                                                                                             Synthetic
                                                                                                                                                                                                                              Synthetic human tropoelastin
                                                                                                                                                                                                                                                                          25-MAR-2003
22-MAR-1995
                   16-DEC-1993;
                                                             07-JUL-1994.
                                                                                                   WO9414958-A1
                                                                                                                                                                                       Tropoelastin;
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                                                                                                                                                                                                                                                                                                                                                                                  AAR56653 standard; protein; 733 AA
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                                                                                                                                                                                                                                                                        (revised)
(first entry)
                                                                                                                                                                                       pharmaceutical; surgical dressing
                   93WO-AU000655
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Pred. No. 4
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RESULT 7
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human synthetic tropoelastin is susceptible to hydrolytic breakdown of the crosslinks. Such material may be useful in e.g. surgical applications, where the gradual loss of material over time is intended. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 733 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 30; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic polynucleotide(s) - encode recombinant tropoelastins and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N-PSDB; AAQ70941.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1994-263633/32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Weiss AS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-DEC-1992;
28-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                         07-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                AAY01301;
                                                                                                                                                                                                                                                                                                                                                                                                     AAY01301 standard;
                                         New derivatives of human tropoelastin - with elastin-like or macromolecular binding properties, useful e.g. as surgical implants.
                                                                                                                                                            18-JUL-1997;
                                                                                                                                                                                  17-JUL-1998;
                                                                                                                                                                                                        28-JAN-1999.
                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                            Tropoelastin; hTE; elastin; glycosaminoglycan; GAG-binding; medical;
                                                                                                                                                                                                                                                                                                                                   Amino acid sequence of synthetic human tropoelastin SHEL.
                    Disclosure; Fig 1; 82pp; English
                                                                            N-PSDB; AAX27704.
                                                                                        WPI; 1999-132162/11.
                                                                                                              Weiss AS
                                                                                                                                                                                                                               WO9903886-A1
                                                                                                                                                                                                                                                                                       hand lotion; surgical implant; industrial product; human; SHEL.
                                                                                                                                                                                                                                                                                                   pharmaceutical; veterinary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UNSY ) UNIV SYDNEY.
                                                                                                                                    (UNSY ) UNIV
                                                                                                                                                                                                                                                    domo sapiens.
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93AU-00009661.
                                                                                                                                                           97AU-00008117.
                                                                                                                                                                                  98WO-AU000564.
                                                                                                                                                                                                                                                                                                                                                                                                     protein;
                                                                                                                                                                                                                                                                                                                                                         entry)
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                                                                                                                                                                                                                                                                                                   cosmetic application; anti-wrinkle;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 501; DB 2;
Pred. No. 4.4e-34;
; Mismatches 0;
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The invention relates to a derivative or variant of human tropoelastin

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RESULT 8
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                                  The present invention relates to the use of an agent that promotes elastin signaling in smooth muscle cells for decreasing or preventing occlusion of a body vessel by smooth muscle cells, decreasing vascular obstruction, promoting actin stress fiber formation or actin polymerisation, increasing F:G actin ratio in a smooth muscle cell, treating or preventing obstructive vascular disease (e.g. restenosis), or preventing stenosis. The agent that promotes elastin signaling in smooth muscle cells is useful for decreasing or preventing occlusion of a body vessel by smooth muscle cells, decreasing vascular obstruction, promoting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Use of an agent that promotes elastin signaling in smooth muscle cells for e.g. decreasing or preventing occlusion of a body vessel by smooth muscle cells, treating or preventing obstructive vascular disease, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; tropoelastin; elastin; body vessel occlusion; stenosis; vascular smooth muscle cell; elastin signaling; vasotropic.
                                                                                                                                                                                                                                                                                                                                                                                                  Claim 24; Page 138-141; Opp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-833516/77.
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No. 4.4e-34;
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The invention relates to a novel method for diagnosing and analysing autoimmune disease or arthritides. The method comprises obtaining a patient sample containing mRNA, analysing gene expression using the mRNA that results in a gene expression signature of the mRNA, and using that gene expression signature to diagnose or analyse the autoimmune disease or arthritides in the patient, where gene expression of at least 60% of the genes correlates with that of the gene signature. The invention further comprises: a treatment of rheumatoid arthritis; identification of genes for targeting in the treatment of rheumatoid arthritis in a mammal, an other than a mouse; diagnosis of rheumatoid arthritis in a mammal, an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         autoimmune disease; arthritide; gene expression analysis; rheumatoid arthritis; collagen-induced; immunosuppressive; antirheumatianthritic; osteopathic; antigout; antiinflammatory; dermatological; immunomodulatory; lupus; ankylosing spondylitis; Fibrositis; fibromyalgia; osteoarthritis; gout; juvenile rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       vascular usecus stenosis. It is also useful in manuractu treatment or prevention of occlusion of treatment or prevention protein as shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          actin ratio in a smooth muscle cell, treating or preventing obstructive vascular disease (e.g. restenosis following angioplasty), or preventing stenosis. It is also useful in manufacturing a medicament for the treatment or prevention of occlusion of a vessel. The present sequence the human tropoelastin protein as shown in the exemplification of the
                                                                                                                                                                                                                                 Diagnosing and analyzing autoimmune disease using gene expression profiles and microarray technology, useful for diagnosing and treating rheumatoid arthritis, lupus, fibrositis, osteoarthritis, fibromyalgia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
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                                                                                                                                                                                 Disclosure; Page;
                                                                                                                                                                                                                                                                                              GENBANK; NP_000492
                                                                                                                                                                                                                                                                                                                WPI; 2003-712740/67.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           501; DB 7;
No. 4.6e-34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 757;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antirheumatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 583
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array or gene chip, specific for rheumatoid arthritis; diagnosis or canalyses of autoimmune disease or rheumatoid arthritis; screening the efficacy of a candidate drug in vitro for the treatment of collagen-induced arthritis; and reducing the symptoms associated with collagencrinduced arthritis. The compositions of the invention have the following activities: immunosuppressive, antirheumatic, antiarthritic, osteopathic, antigout, antiinflammatory, dermatological, and immunomodulatory. The methods and compositions of the present invention are useful for diagnosing and treating autoimmune disease or arthritides, such as fibrownyalgia, osteoarthritis, gout, juvenile rheumatoid arthritis, and an immune disease caused by an infectious agent. This sequence represents of autoimmune diseases or arthritides. Note: This sequence is not shown in the specification. It has been supplied in an electronic format from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 101
(hTE) having elastin-like and/or macromolecule (specifically glycosaminoglycan (GAG))-binding properties. Calls containing vectors comprising the nucleic acids encoding the variants or derivatives are used to produce the proteins recombinantly. The tropoelastin derivatives
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tropoelastin; hTE; elastin; glycosaminoglycan; GAG-binding; medical; pharmaceutical; veterinary; cosmetic application; anti-wrinkle; food; hand lotion; surgical implant; industrial product; human; SHEL; variant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                     New derivatives of human tropoelastin - with elastin-like or macromolecular binding properties, useful e.g. as surgical i
                                                                                                                                                                                                                                                                                                                                                                      17-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human tropoelastin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY01310
                                                                           The invention
                                                                                                                   Claim 35;
                                                                                                                                                                                                               WPI; 1999-132162/11.
                                                                                                                                                                                                                                                                                             (UNSY ) UNIV SYDNEY
                                                                                                                                                                                                                                                                                                                               18-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                          28-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                WO9903886-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY01310 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     584
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       101; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 VAAAAKSAAKVAAKAQLRAAGLGAGIPGLGVGVGVGVPGLGVGAGVPGLGVGAGVPGFGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               VAAAAKSAAKVAAKAQLRAAAGLGAGI PGLGVGVGVPGLGVGAGVPGLGVGAGVPGFGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      757
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADEGVRRSLSPELREGDPSSSQHLPSTPSSPRVPGALAAAK 101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADEGVRRSLSPELREGDPSSSQHLPSTPSSPRVPGALAAAK 624
                                                                                                                 Page 11; 82pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                           relates to a derivative or variant of human
                                                                                                                                                                                                                                                                                                                                   97AU-00008117
                                                                                                                                                                                                                                                                                                                                                                        98WO-AU000564.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          derivative SHEL26-36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 501; DB 7;
Pred. No. 4.6e-34;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 757;
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                                                                           tropoelastin
                                                                                                                                                         implants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 583
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RESULT 11
AAY01305
ID AAY01
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Best Local S
Matches 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  or hybrid proteins containing the derivatives are useful in medical, pharmaceutical, veterinary and cosmetic applications, e.g. as anti-wrinkle or hand lotions, also as surgical implants, foods and industrive products. The hybrid protein have controllable GAG-binding properties, depending on presence or absence of a specific fragment, designated peptide 26A, from hTE. The present sequence represents a human tropoelastin derivative SHELZ6-36
                     The invention relates to a derivative or variant of human tropoelastin (hTE) having elastin-like and/or macromolecule (specifically glycosaminoglycan (GAG))-binding properties. Cells containing vectors comprising the nucleic acids encoding the variants or derivatives are used to produce the proteins recombinantly. The tropoelastin derivatives or hybrid proteins containing the derivatives are useful in medical, pharmaceutical, veterinary and cosmetic applications, e.g. as antiphraducts. The hybrid protein have controllable GAG-binding properties, depending on presence or absence of a specific fragment, designated peptide 26A, from hTE. The present sequence represents a human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 216 AA;
                                                                                                                                                                                                                                    New derivatives of human tropoelastin - with elastin-like or macromolecular binding properties, useful e.g. as surgical it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAY01305 standard;
                                                                                                                                                                                                        Claim
                                                                                                                                                                                                                                                                                    N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                    17-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                28-JAN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAY01305
                                                                                                                                                                                                                                                                                                                                                                                       18-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    pharmaceutical;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Iropoelastin; hTE; elastin;
                                                                                                                                                                                                                                                                                                                                                          VINU ( YSMU)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   lotion;
                                                                                                                                                                                                                                                                                                1999-132162/11.
                                                                                                                                                                                                        15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tropoelastin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          79
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 83;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                    AAX27707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAAGLGAGI PGLGVGVGVPGLGVGAGVPGLGVGAGVPGFGAGADEGVRRSLSPELREGDP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SSSOHLPSTPSSPRVPGALAAAK 101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SSSQHLPSTPSSPRVPGALAAAK 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAAGLGAGI PGLGVGVGVPGLGVGAGVPGLGVGAGVPGFGAGADEGVRRSLSPELREGDP
                                                                                                                                                                                                        Fig 8; 82pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    surgical implant;
                                                                                                                                                                                                                                                                                                                                                           SYDNEY
                                                                                                                                                                                                                                                                                                                                                                                       97AU-00008117.
                                                                                                                                                                                                                                                                                                                                                                                                                    98WO-AU000564
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    veterinary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               derivative SHELgamma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               84.6%;
                                                                                                                                                                                                        English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   glycosaminoglycan; GAG-binding; medical;
cosmetic application; anti-wrinkle; food
nt; industrial product; human; SHEL; vari
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 424;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.2e-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                s, e.g. as anti-
foods and industrial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SHEL; variant.
                                                                                                                                                                                                                                         implants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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tropoelastin

derivative

SHELgamma

SS

Sequence

200

B

Query Match Best Local

Similarity

61.5%; 95.3%;

Length 200

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RESULT 12
ADB64761
ID ADB64
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The invention discloses a polynucleotide comprising a sequence selected CC from 1970 fully defined nucleotide sequences which encode novel CC polypeptides. Also claimed is a polypeptide encoded by the polynucleotide CC or its partial peptide, an antibody binding to the polypeptide or peptide CC of the polynucleotide, immunologically assaying the polypeptide or peptide CC peptide of the polynucleotide by contacting the polypeptide or peptide CC with the antibody of the encoded protein, and observing the binding CC between the two, a transformant carrying the polynucleotide in an CC expressible manner and an antisense polynucleotide. The oligonucleotide is useful as a primer for synthesising the polynucleotide, or as a probe CC for detecting the polynucleotide. The polynucleotide or detecting the polynucleotide. The polynucleotide and encoded CC groteins are useful as pharmaceutical agents and many disease-related CC genes may be included in them, for developing a diagnostic marker or CC medicines for regulation of their expression and activity, or as targets of gene therapy. The genes are involved in tissue and/or cell cregeneration. Membrane proteins, signal transduction-related proteins,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                             Isogai T,
Yamamoto
                                                                                                                                                                                                                                                                                                            New polynucleotides and polypeptides, useful for developing a diagnostic marker or medicines for regulation of their expression and activity, or as targets of gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cell regeneration;
                                                                                                                                                                                                                                                                                  Claim 1;
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25-JAN-2002; 2002US-00350978.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-MAR-2002; 2002EP-00007401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cancer; tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         transcription-related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (HELI-) HELIX RES INST.
(REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                              2003-450961/43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pharmaceutical; diagnostic; gene therapy; tissue regeneration;
egeneration; membrane protein; signal transduction-related prot
ription-related protein; osteoporosis; neurological disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4
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Isono Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       encoded
                                                                                                                                                                                                                                                                              222pp;
                                                                                                                                                                                                                                                                                                                                                                                                                              'n,ĸ
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                                                                                                                                                                                                                                                                                English.
                                                                                                                                                                                                                                                                                                                                                                                                                           Otsuki T, Wakamatsu A,
Hio Y, Otsuka K, Nagai
Otsuka M, Nagahari K, N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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Pred. No. 2.4e-18;
0; Mismatches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NT2RP70003110.
                                                                                                                                                                                                                                                                                                                                                                                                                                Masuho
                                                                                                                                                                                                                                                                                                                                                                                                                 Sato H,
Irie R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ishii S;
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RESULT 13
AAVO1303
ID AAVO1
AC Tropo
KW PAIN
AN WO990
XX PA WO990
XX 18-JU
XX III-JU
XX WPI,
DR N-PSI
XX WPI,
DR N-PSI
XX WPI,
DR N-PSI
XX WPI,
CC The i
CC (hTE)
CC glycc
CC used
CC produ
CC produ
CC produ
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CC produ
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និត្តត្តិត្តត្តិត្តត្តិ
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The invention relates to a derivative or variant of human tropoelastin (hTE) having elastin-like and/or macromolecule (specifically glycosaminoglycan (GAG))-binding properties. Cells containing vectors comprising the nucleic acids encoding the variants or derivatives are used to produce the proteins recombinantly. The tropoelastin derivatives or hybrid proteins containing the derivatives are useful in medical, pharmaceutical, veterinary and cosmetic applications, e.g. as antiwrinkle or hand lotions, also as surgical implants, foods and industrial products. The hybrid protein have controllable GAG-binding properties, depending on presence or absence of a specific fragment, designated peptide 2GA, from hTE. The present sequence represents a human tropoelastin derivative SHELdeltamodified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              encoding them can be used as indicators for diseases (e.g. osteoporos neurological diseases, cancer, tumours. The CDNA may be used to regult the activity or expression of the encoded protein to treat diseases. sequence presented is a protein of the invention. Note: Some of the sequence data for this patent is not represented in the printed specification, but is based on sequence information supplied by the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tropoelastin; hTE; elastin; glycosaminoglycan; GAG-binding; medical; pharmaceutical; veterinary; cosmetic application; anti-wrinkle; food hand lotion; surgical implant; industrial product; human; SHEL.
                                                                                                                                                                                                                                                                                New derivatives of human tropoelastin - with elastin-like macromolecular binding properties, useful e.g. as surgical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
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                                                                                                                                                                                                                                                Claim 7; Fig 3; 82pp; English
                                                                                                                                                                                                                                                                                                                                         N-PSDB; AAX27705.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 VAAAAKSAAKVAAKAQLRAAAGLGAGI PGLGVGVGVFGLGVGAGVPGLGVGAGVPGFGAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                     SYDNEY
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Pred.
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No. 1.4e-17;
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RESULT 14
ADM03792
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Best Local Similarity
                                                                                                                                     The invention relates to a novel human polynucleotide and the encoded polypeptide. A polynucleotide of the invention may have a use in gene therapy. An oligonucleotide of the invention ADM06202-ADM06773 is useful as a primer for synthesizing the polynucleotide or as a probe for detecting the polynucleotide. The polynucleotides ADM01316-ADM03758 are useful in gene therapy, for developing a diagnostic marker or medicines for regulating their expression and activity, or as a target of gene therapy. The proteins ADM03759-ADM06201 encoded by the polynucleotides are useful as pharmaceutical agents. The present sequence represents a sequence represents a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human protein of the invention
                                                                                                                                                                                                                                                                                            New polynucleotides and polypeptides are useful in gene therapy, developing a diagnostic marker or medicines for regulating their expression and activity, or as a target of gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  human;
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                                                                                                   Sequence
                                                                                                                                                                                                                                                                       Claim 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                 Isogai T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-MAR-2002; 2002JP-00137785
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463
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                                                 68; Conserv
                                                                                                                             sequence
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                                                                                                                                                                                                                                                                       SEQ ID NO 2477; 305pp; English.
                                                                                                                                                                                                                                                                                                                                                                                        J, Isono Y,
Yoshikawa T,
                VAAAAKSAAKVAAKAQLRAAAGLGAGIPGLGVGVGVGVGVGAGVPGLGVGAGVPGFGAG
VAAAAKSAAKVAAKAQLRAAAGLGAGIPGLGVGVGVPGLGVGAGVPGLGVGAGVPGFGA-
                                                                                                                                                                                                                                                                                                                                                                                                                 Sugiyama T,
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                                                   Conservative
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                                                            60.8%;
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Hio Y, Otsuka K, Nagai
Otsuka M, Nagahari K, N
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Pred. No. 1.5e-17;
Widematches 0;
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                                                Score 304.5; DB 7;
Pred. No. 1.5e-17;
0; Mismatches 0;
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K, Irie
Masuho Y;
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RESULT 15
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20-AUG-2001; 2001US-0313643P.
20-AUG-2001; 2001US-0313702P.
21-AUG-2001; 2001US-0314031P.
23-AUG-2001; 2001US-0314466P.
28-AUG-2001; 2001US-0315403P.
29-AUG-2001; 2001US-031583P.
17-SEP-2001; 2001US-032394P.
21-SEP-2001; 2001US-032394P.
14-DEC-2001; 2001US-032394P.
14-DEC-2001; 2001US-034033P.
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03-AUG-2001;
07-AUG-2001;
08-AUG-2001;
09-AUG-2001;
13-AUG-2001;
16-AUG-2001;
17-AUG-2001;
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                                                                       Kekuda R, Miller CE, Patturajan M, Pena Shimkets RA, Zerhusen BD, Li I, Ji W, Poss EZ, Boldog FL, Gorman L, Leite MW, Guo X, Zhong M, Gerlach VL, Hjalt T, Ra Edinger SR, Ellerman K, Malyankar UM, Malshorok JP, Lepley DM, Burgess CE, Maj
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19-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-AUG-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         tissue typing; human; NOV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-JAN-2004
                                                                                                                                                                                                                                   19-APR
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                                                                                                                                                                      CURAGEN CORP.
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2002US-0373814P.
2002US-0373825P.
2002US-0373989P.
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2001US-0312892P.
2001US-0313201P.
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2001US-0311292P.
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2002US-00210172
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                                                                           , Pena CEA, Rieger DK;
Ji W, Padigaru M, Casman SJ;
ite MW, Vernet CAM, Anderson D
t T, Rastelli L, Spytek KA;
UM, Macdougall JR, Stone DJ;
CE, Majumder K, Wolenc AR;
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                                                                                                                    Anderson DW;
                                                                                            Stone DJ;
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New NOVX polypeptides and nucleic acids, useful for preventing

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WPI; 2003-663472/62. N-PSDB; ADE40133.

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Search completed: August 10, 2005, 07:33:50 Job time : 120.095 secs
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Best Local Similarity 67.3%;
Matches 68; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a novel NOVX polypeptide. The polypeptide of the invention demonstrates cardiant, antiarteriosclerotic, hypotensive, cytostatic, ancrectic, antidiabetic, immunosuppressive, anti-HIV, neuroprotective, nootropic, antiparkinsonian, antiasthmatic and gynaecological activities and may be useful in diagnosing, treating or preventing NOVX-associated disorders including cardiomyopathy, atherosclerosis, hypertension, cancer, obesity, diabetes, AIDS, multiple sclerosis, graft-versus-host disease, Alzhahmer's disease, Parkinson's disease, asthma or fertility disorders. Furthermore, the polypeptides may be utilised as vaccines whilst the nucleic acids may be used as hybridisation probes, in gene therapy, chromosome mapping, tissue typing, preventive medicine and pharmacogenomics. The current sequence is that of the human NOV protein of the invention.
                                                                                                                                                                                                                                                                                                                                                               Sequence 692 AA;
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                                                                                                                                                                              61
                                                                                                                               ADEGVRRSLSPELREGDPSSSQHLPSTPSSPRVPGALAAAK 101
                                                                                                                                                                                                                                                                     Score 304.5; DB 7; Length 692; Pred. No. 1.6e-17; 0; Mismatches 0; Indels 33;
                                                                                           ----VPGALAAAK 559
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WIS PROFES BURNEY

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Maximum Match 100%
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Maximum DB
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seq length: 2000000000
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Copyright (c) 1993 - 2005 Compugen Ltd
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1011
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   US-08-911-364-1
US-09-340-756E-1
US-09-346-662-1
US-08-464-700-2
US-08-678-039A-70
US-08-397-633A-76
US-08-397-633A-76
US-08-397-633A-68
US-08-397-633A-68
US-08-397-633A-68
US-08-435-641-15
US-08-4707-237A-96
US-08-642-246-15
US-08-642-246-15
US-08-806-029-35
US-08-806-029-35
US-08-806-029-39
US-08-8175-155-59
US-08-175-155-46
US-08-175-155-46
US-08-482-085B-81
US-08-482-085B-81
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US-08-175-155-46
US-08-482-085B-81
US-08-482-085B-81
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Sequence 1, Appli
Sequence 1, Appli
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Sequence 2, Appli
Sequence 40, Appli
Sequence 8, Appli
Sequence 75, Appl
Sequence 54, Appl
Sequence 68, Appl
Sequence 96, Appl
Sequence 15, Appl
Sequence 15, Appl
Sequence 15, Appl
Sequence 15, Appl
Sequence 19, Appl
Sequence 94, Appl
Sequence 52, Appl
Sequence 52, Appl
Sequence 66, Appl
Sequence 66, Appl
Sequence 81, Appl
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29.9	29.9	29.9	29.9	29.9	29.9	29.9	29.9	29.9	29.9	30.1	30.1	30.1	30.5	30.5	30.5	30.5	30.5
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Sequence :	Sequence 2	Sequence	Sequence	Sequence 3	Sequence :	Sequence 2	Sequence 2	Sequence 2	Sequence 1	Sequence :	Sequence	Sequence :	Sequence	Sequence	Sequence	Sequence	Sequence '
25,	25,	25,	103,	36	77,	26,	26,	26,	104,	25,	78,	73	82,	82,	53,	82,	47,
Appl	Appl	Appi	App	Appl	App1	Appl	Appl	App1	App	Appl	Appl	IggA	Appl	Appl	Appl	App1	Appl

ALIGNMENTS

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; STRANDEDNESS:
; TOPOLOGY: lin
; MOLECULE TYPE:
US-08-911-364-1
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                                                                                                                                                                                                                                                     COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/911,364
FILING DATE: 07-AUG-1997
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/023,552
FILING DATE: 07-AUG-1996
ATTORNEY/AGENT INFORMATION:
NAME: Bent, Stephen A.
REGISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 041082/0104
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 672-5300
TELEPHONE: (202) 672-5300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1, Application Patent No. 5969106
        Query Match
Best Local Similarity
Matches 101;
                                                                                                                                                                                                                        TELEFAX: (202) 672-53: INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: ROTHSTEIN, ASET
APPLICANT: ROTHSTEIN, ASET
APPLICANT: KEELY, Fred W.
APPLICANT: ROTHSTEIN, Steven J.
APPLICANT: ROTHSTEIN, Steven J.
APPLICANT: ROTHSTEIN, STEVEN J.
APPLICANT: ROTHSTEIN, STEVEN J.
APPLICANT: ROTHSTEIN AND OTHER FIBROUS PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 20007-5109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 731 amino acid
TYPE: amino acid
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STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 3000 K SI
CITY: Washington
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Conservative
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100.0%; Score 501; DB 2;
100.0%; Pred. No. 2.2e-41;
tive 0; Mismatches 0;
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                                          DB 2;
                                        Length 731;
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RESULT 3
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CURRENT APPLICATION NUMBER: US/09/340,736E
CURRENT FILING DATE: 199-06-29
PRIOR APPLICATION NUMBER: 08/911,364
PRIOR FILING DATE: 1997-08-07
PRIOR APPLICATION NUMBER: 60/023,552
PRIOR FILING DATE: 1996-08-07
NUMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin Ver. 2.1
                                                                                     US-09-964-662-1
                                                                                                                                                            GENERAL INFORMATION:

APPLICANT: PROTEIN SPECIALTIES LTD.

APPLICANT: HSC RESEARCH AND DEVELOPMENT LIMITED PARTNERSHIP

TITLE OF INVENTION: SELF-ALIGNING PEPTIDES MODELED ON HUMAN ELASTIN AND

TITLE OF INVENTION: OTHER FIBROUS PROTEINS

FILE REFERENCE: 041082/0112

CURRENT APPLICATION NUMBER: US/09/964,662

CURRENT FILING DATE: 203-05-08

PRIOR APPLICATION NUMBER: 09/340,736

PRIOR APPLICATION NUMBER: 09/340,736

PRIOR PILING DATE: 1999-06-29

NUMBER OF SEQ ID NOS: 11

SOPTWARE: Patentin Ver. 2.1

SEQ ID NO 1

FENGTH. 731
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1, Application US/09964662 Patent No. 6765086
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Best Local Similarity
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LENGTH: 731
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Patent No. 6489446
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APPLICANT: ROTHSTEIN, ASER
                   Query Match
Best Local Similarity
  Matches 101;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: KEELEY, FRED
APPLICANT: ROTHSTEIN, STEVEN
TITLE OF INVENTION: SELF-ALIGNING PEPTIDES MODELED ON HUMAN ELASTIN
TITLE OF INVENTION: AND OTHER FIBROUS PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                  LENGTH: 731
TYPE: PRT
ORGANISM: Homo sapiens
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  Conservative
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                 100.0%; Score 501; DB 4; Length 731; 100.0%; Pred. No. 2.2e-41;
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                                                                                                                                                                                                                  ; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-464-700-2
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                                                                                                                                              Matches
                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: AU PL6520
FILING DATE: 22-DEC-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: AU PL9661
PILING DATE: 28-UUN-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/AU93/00
PILING DATE: 16-DEC-1993
ATTORNEY/AGENT INFORMATION:
NAME: BEL MACKER

NAME: BEL MACKER

PRIOR APPLICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                            TELEFAX: 215-540-5818
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/464,700
FILING DATE: 7-JUN-1995
                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 733 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: WEISS, ANTHONY S
APPLICANT: MARTIN, STEPHEN L
TITLE OF INVENTION: SYNTHETIC
NUMBER OF SEQUENCES: 54
                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: 215-540-9200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Bak, Mary E.
REGISTRATION NUMBER: 31
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                       TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: Spring House Corporate CITY: Spring House STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE:
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560 ADEGVRRSLSPELREGDPSSSQHLPSTPSSPRVPGALAAAK 600
                                                                      500 VAAAAKSAAKVAAKAQLRAAAGLGAGIPGLGVGVGVPGLGVGAGVPGLGVGAGVPGFGAG
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                  ADEGVRRSLSPELREGDPSSSQHLPSTPSSPRVPGALAAAK 101
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Pred. No. 2.3e-41;
); Mismatches 0;
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RESULT 5 US-08-678-039A-40

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                                                                                                                                                                                                                                                                                                         Sequence 8, Applic Patent No. 6780594
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
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SOFTWARE:
               APPLICANT: WINTERHAGER, ELKE
APPLICANT: REGIDOR, PEDRO
APPLICANT: REGIDOR, PEDRO
APPLICANT: SCOTTI, SIMONE
TITLE OF INVENTION: METHOD FOR IN VITRO DIAGNOSIS
FILE REFERENCE: SCH-1789
CURRENT APPLICATION NUMBER: US/09/961,403
CURRENT FILING DATE: 201-09-25
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Keating, Mark T.
APPLICANT: Morris, Colleen A.
                                                                                                                                                                             APPLICANT: HE-STUMPP, HOLGER
APPLICANT: HAENULER, BERNARD
APPLICANT: KRAETZSCHWAR, JOERN
APPLICANT: KREFT, BERTHOLT
APPLICANT: WINTERHAGER, ELKE
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/678,039A
FILING DATE: 10-UUL-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 792 amino acids
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MEDIUM TYPE: Floppy disk
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
TITLE OF INVENTION:
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NAME: Saxe, Stephen A.
REGISTRATION NUMBER: 38,609
REFERENCE/DOCKET NUMBER: 23:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 555 Inter
STREET: Tower
CITY: Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                      619 ADEGVRRSLSPELREGDPSSSQHLPSTPSSPRVPGALAAAK 659
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555 Thirteenth Street, N.W., Suite 701 East
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        U.S.A.
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Pred. No. 2.4e-41;
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GENERAL INFORMATION:
APPLICANT: Cappello, Joseph
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Best Local S
                                                                                                                             Matches '
                                                                                                                                                          Query Match
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TELEX: 910 277299
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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ORGANISM: Homo sapiens
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PATEN: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/397,633A
FILING DATE:
CLASSIFICATION: 530
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Cappello, Joseph
TITLE OF INVENTION: PRODUCTS COMPRISING SUBSTRATESCAPABLE
TITLE OF INVENTION: OF ENZYMATIC CROSS-LINKING
NUMBER OF SEQUENCES: 105
                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Rowland, Bertram I
REGISTRATION NUMBER: 20,
                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: A-58848-1/BIR PROP-011-1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
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                                                                                                                                                                                                                            TOPOLOGY: 1ii
                                                                                                                                                                                                                                                            TYPE: amino acid
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                                                             68;
 224 PGAGAGSGAGAGSGAGAM----
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Similarity 67.3%;
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                                                                                                                                            Similarity
                                                                                 AAAKSAAKVAAKAQLRAAAGLGAGIPGLGV------GVGVPGLGV-GAGVPGLGV-GAGV 54
                            PGFGAGADEGVRRSLSPELREGDPSSSQHLPSTPSSPRVPGAL 97
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linear
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                                                                                                                                                                                                             protein
                                                                                                                                            31.7%;
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Pred. No. 3.5e-08;
8; Mismatches 40
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Pred. No. 4.8e-22;
0; Mismatches 0;
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 -DPGRYHMAAKGDRAPGTPGGL 262
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                                                                                                                                                                                                                                         Sequence 54, Application US/08397633A Patent No. 5773577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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APPLICANT: Cappel
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INFORMATION FOR SEQ ID NO: 76:
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NAME: ROWLAND, Bertram I
REGISTRATION NUMBER: 20,015
REFERENCE/DOCKET NUMBER: A-:
TELECOMMUNICATION INFORMATION:
                                                                                                                                           APPLICANT: Cappello, Joseph
APPLICANT: Cappello, Joseph
TITLE OF INVENTION: PRODUCTS COMPRISING SUBSTRATESCAPABLE
TITLE OF INVENTION: OF ENZYMATIC CROSS-LINKING
NUMBER OF SEQUENCES: 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 287 amino acids
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MEDIUM TYPE: Floppy disk
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                                                    NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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                                                                     STREET: 4 Luca
CITY: San Francisco
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
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                                                  COUNTRY: USA
                                                                                                                      ADDRESSEE:
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5. 5773577
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                                    94111-4187
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4 Embarcadero Center, Suite 3400
                                                                                             E: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
4 Embarcadero Center, Suite 3400
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PRODUCTS COMPRISING SUBSTRATESCAPABLE
OF ENZYMATIC CROSS-LINKING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 158; DB 1; Length 287; Pred. No. 4.5e-08;
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APPLICALL.

PILING DATE:

FILING DATE:

CLASSIFICATION: 530

ATTORNEY/AGENT INFORMATION:

ATTORNEY/AGENT INFORMATION:

NAME: ROWland, Bertram I

REGISTRATION NUMBER: 20,015

REFERENCE/DOCKET NUMBER: A-58848-1/BIR PROP-011-1

FELECOMMUNICATION INFORMATION:

TELECHONE: (415) 781-1989

TELECHONE: (415) 398-3249

TELECHONE: (415) 398-3249

TELECHONE: (415) 398-3249
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APPLICANT: Cappel
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Cappello, J
TITLE OF INVENTION: PR
TITLE OF INVENTION: OF
NUMBER OF SEQUENCES: 1
                                                                                                                                                                                                                                         COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/397,633A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 877 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 781-1989
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPAUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Pelear- ""
DESERVE - ""
                                                                                                                                                                                                                                                                                                                                                                                                          CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                         ZIP:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Rowland, Bertram REGISTRATION NUMBER: 20
                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (415) 781-198
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRODUCTS COMPRISING SUBSTRATESCAPABLE OF ENZYMATIC CROSS-LINKING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      105
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Center, Suite 3400
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Pred. No. 2.9e-07;
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; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-397-633A-68
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GENERAL INFORMATION:
APPLICANT: Stedro
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                                                                                                                                                             Query Match
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, vc

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/435,641

FILING DATE: 05-MAY-1995

CLASSIFICATION: 514

ATTORNEY/AGENT INFORMATION:

NAME: Royland Retrem I
                                                                                                                                                                                                                                                                                          TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                          NAME: Rowland, Bertram I.
REGISTRATION NUMBER: 20015
REFERENCE/DOCKET NUMBER: A-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
                                                                                                                                                                                                             MOLECULE TYPE: peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         / Match 30.7%;
Local Similarity 45.7%;
les 42; Conservative
                                                                                                                                              Local
                                                                                                                                                                                                                              STRANDEDNESS:
TOPOLOGY: li
                                                                                                                                                                                                                                                             TYPE: amino acid
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                                                                                                                                                                                                                                                                              ENGTH:
851
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                                                                                                                                            Similarity
                                                           PGAGAGSGAGAGSGAGAM----DPGRYQDLRS 878
                                                                                            AAAKSAAKVAAKAQLRAAAGLGAGIPGLGV-GVGVPGLGV-----GAGVPGLGV-GAGV 54
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                                                                                                                                                                                                                                                                              884 amino acids
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                                                                                                                              Conservative
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                                                                                                                                                                                                                                            single
                                                                                                                                          30.7%;
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                                                                                                                             Score 154; DB 2; Length 884; Pred. No. 3.7e-07; Mismatches 31; Indels
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Suite 3400
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US-08-707-237A-96
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APPLICATION DATA:

PRIOR APPLICATION UMBER: US 08/053,04>

APPLICATION NUMBER: US 08/053,04>

FILING DATE: 22-APR.1993

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/609,716

APPLICATION NUMBER: US 07/609,716

APPLICATION NUMBER: US 07/609,716

APPLICATION NUMBER: US 07/609,716
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                                                                                                                            Matches
                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                   TELEX: 910 277299
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PILING DATE: 29-OCT-1987
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 06/
FILING DATE: 04-NOV-1986
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/175,155
FILING DATE: 29-DEC-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 09-NOV-PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                    MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
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                                                                                                                                            Local
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REFERENCE/DOCKET NUMBER: A-55186-10/WHD
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                                                                                                                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                       TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                               TELEFAX:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                      LENGTH:
                                                    55 PGFGAGADEGVRRSLSPELREGDPSSSQHLPS 86
                                                                                      3 AAAKSAAKVAAKAQIRAAAGIGAGIPGIGV-GVGVPGIGV-----GAGVPGIGV-GAGV 54
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                                                                                                                                            Similarity
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Capello, Joseph
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                                                                                                                                                                                                                                     unknown
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29-OCT-1987
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                                                                                                                                                                                                                                                      unknown
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                                                                                                                        Score 154; DB 2; I
Pred. No. 3.7e-07;
7; Mismatches 31;
                                                                                                                                                            Length 884;
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RESULT 13
US-08-642-246-15
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US-09-451-206-15
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Best Local S
Matches 42
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Patent No. 6423333
GENERAL INFORMATION:
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APPLICANT: STEDRONSKY, Erwin R.
APPLICANT: CAPPELLO, Joseph
TITLE OF INVENTION: Tissue Adhesive
TITLE OF INVENTION: Crosslinking
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 415-398-3249
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 884 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: A6
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/642,246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT STREET: Four Embarcadero Center, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity les 42; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: ROWLAND, Bertram I REGISTRATION NUMBER: 20015
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                  CAPPELLO, Joseph
TITLE OF INVENTION: Tissue Adhesive Using Synthetic
                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: FLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
                                                                                  NUMBER OF SEQUENCES: 35
                                                                                                                                                   APPLICANT: STEDRONSKY, Erwin R.
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                                                                                                                                                                                                                                                                                                                  PGFGAGADEGVRRSLSPELREGDPSSSOHLPS 86
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STATE: CA
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                                                                                                   Crosslinking
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Pred. No. 3.7e-07;
7; Mismatches 31;
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                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                         ZIP:
                                                                                                                                                                                                                                     COUNTRY:
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Sequence 15, Application PC/TUS9606229 GENERAL INFORMATION:
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                CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: ROWLAND, Bertram I
REGISTRATION NUMBER: 200
                                                                                                                               COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Tissue Adhes: TITLE OF INVENTION: Crosslinking
                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS: PLEHR, HOHBACH, TEST,
                                                                                                                                                                                                                                                                                        STREET: rour
CITY: San Francisco
                                                                                          APPLICATION NUMBER: FILING DATE:
REFERENCE/DOCKET NUMBER: A61127-1/BIR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible operating system: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, V CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/451,206
FILING DATE: 29-No. 6423333-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/642,246
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              55 PGFGAGADEGVRRSLSPELREGDPSSSQHLPS 86
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 884 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: 415-781-1989
TELEFAX: 415-398-3249
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                                                                                                                                                                                                                                                                                                                             Four Embarcadero Center,
                                                                                                                                                                                                                                                                                                                                                                                                                            STEDRONSKY, Erwin R.
CAPPELLO, Joseph
WENTION: Tissue Adhesive Using Synthetic
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                                                                                                                                                                                                            Floppy disk
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                  20015
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Pred. No. 3.7e-07;
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Suite 200
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 s
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Maximum DB seq length: 200000000
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Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Published Applications AA:*
|: /cgn2_6/ptodata/1/pubpaa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.
                        (cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
(cgn2_6/ptodata/1/pubpaa/US11_NEW_PUB.pep:*
(cgn2_6/ptodata/1/pubpaa/US11_NEW_PUB.pep:*
(cgn2_6/ptodata/1/pubpaa/US60_NEW_PUB.pep:*
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jn2 6/ptcodata/1/pubpaa/US10C_PUBCOMB.pep:*
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jn2 6/ptcodata/1/pubpaa/US10E_PUBCOMB.pep:*
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2 6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
2 6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
2 6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
2 6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*
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PUBCOMB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

### SUMMARIES

1 2 2 3 4 4 4 7 7 10 11	Result No.
501 490 304.5 304.5 304.5 304.5 304.5 304.5 301.5	Score
100.0 97.8 97.8 60.8 60.8 60.8 60.8 60.8 30.7	Query Match
731 788 788 617 663 692 730 730 731 711 244	Query Match Length DB
16 15 15 15 16 16	:
US-09-964-662-1 US-10-852-065-4 US-10-852-575-3 US-10-104-047-2915 US-10-108-260A-2477 US-10-210-172-40 US-09-961-403-8 US-10-723-860-2566 US-10-210-172-38 US-10-210-172-38 US-10-845-775A-27	ID
Sequence 1, Appli Sequence 4, Appli Sequence 2915, Ap Sequence 2917, Ap Sequence 2477, Appl Sequence 40, Appli Sequence 8, Appli Sequence 38, Appl Sequence 38, Appl Sequence 37, Appl Sequence 27, Appl	Description

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7-931-	306-029-2	-10-096-986-	-10-096-98	-10-939-036-	-10-845-936A	-10-845-	-10-800-179-3	-10-096-986-9	-10-939-0	-10-845-936	-10-845-775A-3	-10-800-179-3	08-806-029-19	US-10-939-036-25	-10-845-936A-	-10-845-775A-	179-2	08-806-0	-10	-10-845-936A-3	-10-845-	-10-845-936A-	-10-117-931-15	US-10-939-036-28	-10-939-036-	0-845-	0-845-936A-2	45-775A-	-10-845-775A-2	US-10-800-179-28	-800-179-2	US-10-939-036-27	US-10-845-936A-27	
Sequence 26, Appl	25,	82,	e 81,	e 30,	e 30,	æ	e 30,	æ	31,	31,	3 L	)   	19	e 25	25	25	e 25	35, App1	Sequence 34, Appl	ω w	32	37,		28,	26,	28,	26,	82.	26,	28	26,	Ø	equence 27,	

## ALIGNMENTS

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NUMBER OF SEQ ID NOS: 11
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 731
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-964-662-1
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APPLICANT: PROTEIN SPECIALTIES LTD.
APPLICANT: HSC RESEARCH AND DEVELO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1, Application US/09964662
Publication No. US20030166846A1
                                                                                                                                                                   Matches
                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: HSC RESEARCH AND DEVELOPMENT LIMITED PARTNERSHIP TITLE OF INVENTION: SELF-ALIGNING PEPTIDES MODELED ON HUMAN I TITLE OF INVENTION: OTHER FIBROUS PROTEINS FILE REFERENCE: 041082/0112
                                                                                                                                                                                                                                                                                                                                                                                           CURRENT FILING DATE: 2003-05-08
PRIOR APPLICATION NUMBER: 09/340,736
PRIOR FILING DATE: 1999-06-29
                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/964,662
CURRENT FILING DATE: 2003-05-08
                                                                                498 VAAAAKSAAKVAAKAQLRAAAGLGAGIPGLGVGVGVPGLGVGAGVPGLGVGAGVPGFGAG
558 ADEGVERSISPELREGDPSSSQHLPSTPSSPRVPGALAAAK 598
                                                                                                                                                                   101;
                          61 ADEGVRRSLSPELREGDPSSSQHLPSTPSSPRVPGALAAAK 101
                                                                                                             1 VAAAAKSAAKVAAKAQIRAAAGIGAGIPGIGVGVGVGVEGIGVGAGVPGIGVGAGVPGFGAG 60
                                                                                                                                                               100.0%; Score 501; DB 10; ilarity 100.0%; Pred. No. 2.5e-35; Conservative 0; Mismatches 0;
                                                                                                                                                                                                          Length 731;
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; ORGANISM: Homo sapiens US-10-852-575-3
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APPLICANT: CENIZO, Val, rie
APPLICANT: BOUEZ, Charbel
APPLICANT: SOMMER, Pascal
APPLICANT: DAMOUR, Odile
APPLICANT: GLEYZAL, Claudine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: REYMERMIER, Corinne
TITLE OF INVENTION: Stimulation of the synthesis and of the activity of an isoform
TITLE OF INVENTION: of lysyl oxidase-like LOXL for stimulating the formation of
TITLE OF INVENTION: elastic fibers
FILE REFERENCE: 11123.81US01
CURRENT FILING NUMBER: US/10/852,065
CURRENT FILING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: FR 03 07177
PRIOR FILING DATE: 2003-06-13
NUMBER OF SEQ ID NOS: 31
SOFTWARE: PatentIn version 3.1
SEQ ID NO 4
LENGTH: 788
                                      NUMBER OF SEQ ID NOS: 31
SOPTWARE: Patentin version 3.1
SEQ ID NO 3
LENGTH: 788
TYPE: PRT
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Publication No. US20040253220A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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Publication No. US20040258676A1
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                                                                                                                                                   APPLICANT: REYMERMIER, Corinne
APPLICANT: REYMERMIER, Corinne
APPLICANT: ORLY, Isabelle
TITLE OF INVENTION: Stimulation of the activity of an isoform of lysyl oxidase
TITLE OF INVENTION: for combating against some pathologies due to an incomplete,
TITLE OF INVENTION: absent or disorganized elastogenesis
FILE REFERENCE: 11123 82USO1
CURRENT APPLICATION NUMBER: US/10/852,575
CURRENT FILING DATE: 2004-05-24
PRIOR APPLICATION NUMBER: FR 03 07178
PRIOR PILING DATE: 2003-06-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: PERRIER, Bric
APPLICANT: BOUEZ, Charbel
APPLICANT: CENIZO, Valerie
APPLICANT: SOMMER, Pascal
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                                                                                                                                                                                                                                                                                                                                                                     DAME, CILL GLEYZAL, CILL COP
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GLEYZAL, Claudine
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ANDRE, Val, rie
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98.1%;
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Pred. No. 2.4e-34;
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US-10-104-047-2915
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CURRENT FILING DATE: 2002-03-27
NUMBER OF SEQ ID NOS: 5458
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2477
LENGTH: 663
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2915, Application US/10104047 Publication No. US20030236392A1 GENERAL INFORMATION:
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SOPTWARE: PatentIn Ver. 2.1
SEQ ID NO 2915
LENGTH: 617
TYPE: PRT
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Best Local Similarity
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Best Local Similarity
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                                                                                                                                   Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                               APPLICANT: HELIX RESEARCH INSTITUTE
TITLE OF INVENTION: No. US20040005560Alel full length cDNA
FILE REFERENCE: H1-A0106
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                                                                                                                 68;
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61 ADEGVRRSLSPELREGDPSSSQHLPSTPSSPRVPGALAAAK 101
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                                                                         1 VAAAAKSAAKVAAKAQLRAAAGLGAGIPGLGVGVGVPGLGVGAGVPGLGVGAGVPGFGAG
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98.1%;
                                                                                                                                 60.8%; Score 304.5; DB 15; Length 663; 67.3%; Pred. No. 2.7e-18;
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Pred. No. 2.5e-18;
0; Mismatches 0
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Pred. No. 2.4e-34;
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                                    ; LENGTH: 692
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-210-172-40
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PRIOR FILING DATE: 2001-08-02
PRIOR APPLICATION NUMBER: 60/323,994
PRIOR FILING DATE: 2001-09-21
PRIOR APPLICATION NUMBER: 60/373,814
PRIOR FILING DATE: 2002-04-19
PRIOR APPLICATION NUMBER: 60/310,291
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Query Match
                                                                                                        SOFTWARE: CuraSeqList version 0.1 SEQ ID NO 40
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                                                                                                                                            Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 327
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CURRENT FILING DATE: 2001-08-01
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APPLICANT: Alsobrook II, John
APPLICANT: Lepley, Denise et al.
TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NUCLEIC ACIDS ENCODING SAME, AND METHOI
                                                                                                                                                                                  PRIOR FILING DATE: 2001-08-16
                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/312,892
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PRIOR APPLICATION NUMBER: 60/310,544
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                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 60/311,292 FILING DATE: 2001-08-09
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Spytek, Kimberly
Edinger, Shlomit
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Malyankar, Uriel
MacDougall, John
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Leite, Mario
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Casman, Stacie
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Shimkets, Richard
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Pena, Carol
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60.8%;
Score 304.5;
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DB 15;
Length 692;
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; LENGTH: 730
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-723-860-2566
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US-10-723-860-2566
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SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 8
                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2566, Application US/10723860 Publication No. US20040253606A1 GENERAL INFORMATION:
                                                                                                                           SOFTWARE: PatentIn version 3.2 SEQ ID NO 2566
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Best Local S
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Query Match
                                                                                                                                                                                     APPLICANT: Aziz, Natasha
APPLICANT: Ginsburg, Wendy M.
APPLICANT: Zlotnik, Albert
TITLE OF INVENTION: Methods of Diagnosis of Soft Tissue
TITLE OF INVENTION: Methods for Screening for Soft Tissue
FILE REFERENCE: 05882.0193.NPUS01
CURRENT APPLICATION NUMBER: US/10/723,860
CURRENT FILING DATE: 2003-11-26
PRIOR APPLICATION NUMBER: 60/429,739
PRIOR FILING DATE: 2002-11-26
PRIOR FILING DATE: 2002-11-26
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CURRENT APPLICATION NUMBER: US/09/961,403
CURRENT FILING DATE: 2001-09-25
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APPLICANT: HAENDLER, BERNARI
APPLICANT: KRAETZSCHMAR, JO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
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NUMBER OF SEQ ID NOS: 8393
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APPLICANT: SCOTTI, SIMONE
TITLE OF INVENTION: METHOD FOR IN VITRO DIAGNOSIS OF ENDOMETRIOSIS
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TYPE: PRT
ORGANISM: Homo sapiens
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Local Similarity 67.3%;
es 68; Conservative
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KRAETZSCHMAR, JOERN
KREFT, BERTHOLT
WINTERHAGER, ELKE
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  Score 304.5;
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    DB 16;
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    Length 730;
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                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/310,291
PRIOR FILING DATE: 2001-08-03
PRIOR APPLICATION NUMBER: 60/310,951
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Publication No.
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                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/309,501
                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/10/210,172
CURRENT FILING DATE: 2001-08-01
                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Lepley, Denise et al.
TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NUCLEIC ACIDS ENCODING SAME, AND METHODE REFERENCE: 21402-416 A
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                           PRIOR FILING DATE: 2001-08-16
                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: 60/373,814
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                                           PRIOR APPLICATION NUMBER: 60/312,892
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                                                                                                                                                                  FILING DATE: 2001-08-08
APPLICATION NUMBER: 60/310,544
                                                                                                                                                                                                                                              FILING DATE:
                                                                        APPLICATION NUMBER: 60/313,201
                                                                                                                                     APPLICATION NUMBER: 60/311,292
                                                                                                                                                       FILING DATE:
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                                                                                                      APPLICATION NUMBER: 60/311,979
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Prior Application SEQ ID NOS: 327
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Leite, Mario
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Zerhusen, Bryan
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Spytek, Kimberly
Edinger, Shlomit
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MacDougall, John
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Linda
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              See
              File
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         APPLICANT: Kumar, Manoj
APPLICANT: Kumar, Manoj
APPLICANT: Mazeaud, Isabelle
APPLICANT: Christiano, Steven P.
APPLICANT: Christiano, Steven P.
TITLE OF INVENTION: Controlled Release of Active Age
TITLE OF INVENTION: Protein Polymers
FILE REFERENCE: DOC 0077 PA/GC 792-6/DC 5110
CURRENT APPLICATION NUMBER: US/10/845,775A
CURRENT FILING DATE: 2004-05-14
PRIOR APPLICATION NUMBER: US 60/470,465
PRIOR FILING DATE: 2003-05-14
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APPLICANT: Cuevas, William A.

APPLICANT: Kumar, Manoj

TITLE OF INVENTION: Use of Repeat Sequence Protein

TITLE OF INVENTION: Compositions

FILE REFERENCE: DOC 0057 PA / GC792-4 / DC 5074

CURRENT APPLICATION NUMBER: US/10/800,179

CURRENT FILING DATE: 2004-03-12

CURRENT FILING DATE: 2004-03-12
                                                                                                                                                                     Sequence 27, Application US/10845775A Publication No. US20040228913A1 GENERAL INFORMATION:
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PRIOR FILING DATE: 2003-03-12
NUMBER OF SEQ ID NOS: 31
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TYPE: PRT
ORGANISM: Unknown
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ORGANISM: Homo
NUMBER OF SEQ ID NOS: 31
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Pred. No. 1.2e-05;
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Pred. No. 5.3e-18;
1; Mismatches 0
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                                                                                                             Agents Utilizing Repeat Sequence
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US-10-845-936A-27
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                                                                                                                                                                                                                                                     RESULT 13
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Matches
                                                                                                                                                                Sequence 27, Application US/10939036 Publication No. US20050142094A1 GENERAL INFORMATION:
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SOFTWARE: PatentIn version 3.2
SEQ ID NO 27
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Best Local (
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LENGTH: 244
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APPLICANT: Collier, Katherine D.
APPLICANT: Cuevas, William A.
APPLICANT: Kumar, Manoj A.
APPLICANT: Kumar, Manoj A.
APPLICANT: Kumar, Manoj A.
TITLE OF INVENTION: Repeat Sequence Protein Polymer Active Agent Conjugates, Methods
TITLE OF INVENTION: and Uses
FILE REFERENCE: DOC0068PA/DC5058/GC792
CURRENT APPLICATION NUMBER: US/10/485,936A
CURRENT FILING DATE: 2003-05-14
PRIOR APPLICATION NUMBER: 60/470,464
PRIOR APPLICATION NUMBER: 60/470,464
PRIOR FILING DATE: 2003-05-14
APPLICANT: Kumar, Manoj
TITLE OF INVENTION: Use of Repeat Sequence Protein
TITLE OF INVENTION: in Personal Care Compositions
FILE REFERENCE: GC792-4-2
CURRENT APPLICATION NUMBER: US/10/939,036
CURRENT FILING DATE: 2004-09-10
PRIOR APPLICATION NUMBER: US 10/800,179
PRIOR FILING DATE: 2004-03-12
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ORGANISM: unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 244
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45.7%;
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Pred. No. 1.2e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 154; DB 16;
Pred. No. 1.2e-05;
7; Mismatches 31;
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US-10-939-036-27
                                                                                                                                                    US-10-800-179-28
                                                                                                                                                                        RESULT 15
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        Sequence 28, Application US/10800179
Publication No. US20040180027A1
GENERAL INFORMATION:
APPLICANT: Cuevas, William A.
APPLICANT: Kumar, Manoj
TITLE OF INVENTION: Use of Repeat Sequence Protein Polymers
FILE REFERENCE: DOC 0057 PA / GC792-4 / DC 5074
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Cuevas, William A.
APPLICANT: Kumar, Manoj
TITLE OF INVENTION: Use of Repeat Sequence Protein Polymers
TITLE OF INVENTION: Compositions
TITLE OF INVENTION: Compositions
TILE OF INVENTION: Compositions
TILE REFERENCE: DOC 0057 PA / GC792-4 / DC 5074
CURRENT APPLICATION NUMBER: US/08-00,179
CURRENT APPLICATION NUMBER: US/08-03-12
PRIOR APPLICATION NUMBER: 60/454,077
PRIOR FILING DATE: 2003-03-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn version 3.2
SEQ ID NO 26
LENGTH: 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 27
LENGTH: 244
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 31
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Unknown
FEATURE:
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Similarity 45.7%;
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Compositions
0057 PA / GC792-4 / DC
NUMBER: US/10/800,179
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Pred. No. 1.2e-05;
7; Mismatches 31;
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Pred. No. 1.2e-05;
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Page 6
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Num on:	OHLPSTPSSPRVPGALAAAK 101  QHLPSTPSSPRVPGALAAAK 101  QHLPSTPSSPRVPGALAAAK 101  QHLPSTPSSPRVPGALAAAK 101  Description  Proposable aminorus  Apporthetical prote  GLB1 protein - tom  probable aminorus  hypothetical prote  prote hypothetical prote  probable disease r  glycine-rich prote  probable disease r  glycine-rich prote  protectical prote  probable disease r  glycine-rich prote  prote  prote hypothetical prote
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on: August 10, 2005, 07:25:52; Search	.6
(withou 394.685	
US-09-743-818A-4_COPY_500	
1 VAAAAKSAAKVAAKAQLRAA	SPRVPGALAAAK 101
table: BLOSUM62 Gapop 10.0 , Gapext 0.	
earched: 283416 segs, 96216763	
number of hits satisfying chosen parameters: 2	283416
DB seg length: 0 DB seg length: 2	
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No. is the number of results predicted b greater than or equal to the score of th derived by analysis of the total score	G E
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t Query Score Match Length DB	Description
1 501 100.0 792 1 2 265 52.9 860 1	precurso
264.5 52.8 747 1	precur
231 46.1 770 2	stin -
154.5 30.8 784 2	
130 25.9 907 2	sporozoite surface
117 23.4 1596 2 112.5 22.5 127 2	neurogenic locus m
109.5 21.9 505 2	י ל
109 21.8 532 2	뭐유
104 20.8 151 2	glycine-rich prote
103.5 20.7 108 2	hypothetical prote
103 20.6 100 2	glycine-rich prote
102.5 20.5 129 2	glycine-rich prote
99.5 19.9 130 2	hypothetical prote
99.5 19.9 1901 2	hypothetical glyci fibroin - silkworm
99 19.8 100 1	probable disease r
98.5 19.7 158 2 98.5 19.7 1240 2	prot
98 19.6 143 2 98 19.6 1147 1	chai
97.5 19.5 1733 1 97.5 19.5 1958 2	

94	94.5	94.5	94.5	94.5	95	95	95	95	95	95.5	96	96.5	96.5	96.5	96.5
18.8	18.9	18.9	18.9	18.9	19.0	19.0	19.0	19.0	19.0	19.1	19.2	19.3	19.3	19.3	19.3
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S23463	AC1064	T35317	T43959	S56703	T29351	T29350	A55267	T09084	B96506	E86179	C85356	A35912	P2WL41	T18758	B85356
cuticular protein	outer membrane tim	ര	ical pro	-	collagen alpha 2(I	ğ	collagen alpha 5(I	phosphatidylinosit	hypothetical prote		glycine-rich prote	nomeotic protein o	L2 protein - human	hypothetical prote	_

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hypothetical prote
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A;Note: this sequence represents a composite of several splice forms
R;Fazio, M.J.; Olsen, D.R.; Kuivaniemi, H.; Chu, M.L.; Davidson, J.M.; Rosenbloom, J.;
Lab. Invest. 58, 270-277, 1988
A;Title: Isolation and characterization of human elastin cDNAs, and age-associated variab; Reference number: A53891; MUID:88156138; PMID:2831431
A;Accession: A53891; MUID:88156138; PMID:2831431
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Species: Homo sapiens (man)
C;Date: 22-Un-1990 #sequence revision 26-Jul-1996 #text_change 09-Jul-2004
C;Date: 22-Un-1990 #sequence revision 26-Jul-1996 #text_change 09-Jul-2004
C;Accession: A32707; A33705, A30524; A53891
R;Indik, Z.; Yeh, H.; Ornstein-Goldstein, N.; Sheppard, P.; Anderson, N.; Rosenbloom, J
Proc. Natl. Acad. Sci. U.S.A. 84, 5680-5684, 1987
A;Title: Alternative splicing of human elastin meNA indicated by sequence analysis of c
A;Reference number: A32707; MUID:87289668; PMID:3039501
A;Accession: A32707
                                      A;Map position: 7q11.23-7q11.23
C;Superfamily: elastin
C;Superfamily: elastin
C;Keywords: alternative splicing; extracellular matrix; glycoprotein; hydroxylysine
E;1-26/Domain: signal sequence #status predicted <SIG>
F;27-792/Product: elastin #status predicted <MAT>
F;782-787/Disulfide bonds: #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: GB:J04821; NID:g182052; PIDN:AAA52379.1; PID:g553276 R;Fazio, M.J.; Olsen, D.R.; Kauh, E.A.; Baldwin, C.T.; Indik, Z.; Ornstein-Goldstein, N J. Invest. Dermatol. 91, 458-464, 1988 A;Title: Cloning of full-length elastin cDNAs from a human skin fibroblast recombinant A;Reference number: A30524; MUID:89009960; PMID:3171221 A;Accession: A30524
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Residues: 1-500,507-792 <IND>
A;Cross-references: UNIPROT:P15502; UNIPROT:Q9UMK5; GB:M16983; GB:J02948
A;Cross-references: UNIPROT:P15502; UNIPROT:Q9UMK5; GB:M16983; GB:J02948
A;Cross-references: UNIPROT:P15502; UNIPROT:Q9UMK5; GB:M16983; GB:J02948
A;Cross-reference: UNIPROT:P15502; UNIPROT:Q9UMK5; GB:M16983; GB:J02948
A;Cross-reference:P1500, Complete human elastin gene. Delineation of unusual felareference number: A33705; MUID:89255358; PMID:2722804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       elastin precursor, long splice form - human N;Alternate names: tropoelastin
                                                                                                                                                                                                                                                                                                                                                                                                           A;Molecule type: mRNA
A;Residues: 164-453,483-500,507-617,651-792 <FA2>
A;Cross-references: GB:M24782; NID:g182063; PIDN:AAA53190.1; PID:g182064
C;Comment: The term tropoelastin refers to a soluble precursor form of the extracellula
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Molecule type: mRNA
A; Residues: 1-453,483-617,651-792 <FAZ>
                                                                                                                                                                                                                              A;Cross-references: GDB:119107; OMIM:130160
A;Map position: 7q11.23-7q11.23
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A;Residues: 1-27 <BAS>
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A;Title: Use of an intron length polymorphism to localize the t A;Reference number: A55721; MUID:95130069; PMID:7829060 A;Reference number: A55721 A;MOJecule type: mRNA A;Residues: 1-860 < MYD>
A;Cross-references: UNIPROT:P54320; GB:U08210; NID:g473273; PID A;Cross-references: UNIPROT:P54320; GB:U08210; NID:g473273; PID C;Genetics: A;Map position: 5
A;Map position: 5
C;Superfamily: elastin
C;Keywords: alternative splicing; extracellular matrix; glycopr F;1-27/Domain: signal sequence #status predicted <SIG>F;1-27/Domain: signal sequence #status predicted <MAT>F;850-855/Disulfide bonds: #status predicted
                                                                                                                                                                                                                                                                                                                       N;Alternate names: tropoelastin
N;Contains: elastin precursor, splice form b; elastin precursor, splice form c
C;Species: Bos primigenius taurus (cattle)
C;Date: 08-Jun-1989 #sequence revision 26-Jul-1996 #text change 09-Jul-2004
C;Accession: A31865; A26728; B26728; C26728; A22343; I45886
C;Accession: A31865, A26728; B26728; C26728; A22343; I45886
C;Accession: A31865, A26728; C26728; A22343; I45886
C;Accession: A31865, A26728; B26728; C26728; A22343; I45886
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C;Species: Mus musculus (house mouse)
C;Date: 18-Aug-1995 #sequence_revision 16-Aug-1996 #text_change 09-Jul-2004
C;Accession: A55721
A;Molecule type: mRNA
A;Residues: 1,'RS',4-11,'E',13-636,'V',638-747 <RAJ>
A;Cross-references: GB:J02717; NID:g163019; PIDN:AAA30503.1;
                                                                                                                                                                                                                                                              Biochemistry 28, 2365-2370, 1989
A;Title: Structure of the bovine elastin gene and S1 nuclease
A;Reference number: A31865; MUID:89274159; PMID:2543440
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                                                                                      A; Title: Primary structures of bovine elastin a, b, and A; Reference number: A92640; MUID:87194772; PMID:3032943
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                                                                                                                                                                                                   A; Molecule type: DNA
A; Residues: 1-27 < YEH>
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                                                                                                                                 Chem. 262, 5755-5762,
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Pred. No. 4.4e-14;
3; Mismatches 14
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  PID:g163020
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A;Molecule type: DNA
A;Molecule type: DNA
A;Rossidues: 678-683,685-747 <ROS>
A;Residues: 678-683,685-747 <ROS>
A;Residues: 678-683,685-747 <ROS>
A;Residues: 678-683,685-747 <ROS>
B;Cross-references: GB:M31898; NID:g163015; PIDN:AAA96417.1; PID:g163018
A;Brown, P.L.; Mecham, L.; Tisdale, C.; Mecham, R.P.
Biochem. Biophys. Res. Commun. 186, 549-555, 1992
A;Title: The cysteine residues in the carboxy terminal domain of tropoela
A;Reference number: A58621; MUID:92337651; PMID:1632791
A;Contents: annotation, disulfide bonds
C;Comment: The term tropoelastin refers to a soluble precursor form of t)
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R;Rosenbloom, J.
Lab. Invest. 51, 605-623, 1984
Lab. Invest. 51, 605-623, 1984
-- ^^^coacase. PMID:6150137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Residues: 1,'RS',4-11,'E',13-225,260-636,'V',638-747 <RA3>
A;Residues: 1,'RS',4-11,'E',13-225,260-636,'V',638-747 <RA3>
A;Cross-references: GB:K03506; NID:g163027; PIDN:AAA30506.1; PID:g163028
R;Cicila, G.; May, M.; Ornstein-Goldstein, N.; Indik, Z.; Morrow, S.; Yeb
Biochemistry 24, 3075-3080, 1985
Biochemistry 24, 3075-3080, 1985
A;Title: Structure of the 3' portion of the bovine elastin gene.
A;Reference number: A22343; MUID:85280426; PMID:2992576
                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Keywords: alternative splicing; extracellular matrix; glycoprotein; hydrox f;1-747/Product: elastin precursor, splice form a #status predicted <EPA> F;1-725,260-747/Product: elastin precursor, splice form c #status predicted F;1-225,240-747/Product: elastin precursor, splice form b #status predicted F;1-225,240-747/Product: elastin precursor splice form b #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A,Title: Biology of disease: Elastin: Relation of pro A;Reference number: I45885; MUID:85059254; PMID:61501 A;Accession: I45886
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A; Molecule type:
A; Residues: 1, 'R
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F;27-747/Product: elastin #status predicted <MAT>
F;105,109,252,271,275,324,327,400,404,407,445,448,489,493,544,548,552,606,609,645,649,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Introns: 634/3; 653/3; 676/3; 689/3; 70:
A; Note: the list of introns is incomplete
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A; Residues: 1, 'RS', 4-
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A;Cross-references: GB:K03505; NID:g163025; PIDN:AAA30505.1; PID:g163026
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A;Residues: 613-747
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                                                                                                                                                                                                                                                                                                                                                  F;737-742/Disulfide bonds: #status experimental
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                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                             Local
                                                61 ADEGVRRSLSPELREGDPSSSQHLPSTPSSPRVPGALAAAK 101
                                                                                                                                               60;
                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                      52.8%;
                                                                                                                                                                                                                                 1; Mismatches
                                                                                                                                                                                                                        Score 264.5; DB 1;
Pred. No. 4.2e-14;
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elastin precursor - rat
N;Alternate names: tropoelastin
C;Species: Rattus norvegicus (Norway rat)
C;Date: 11-Jan-1991 #sequence revision 16-Aug-1996 #text_change 09-Jul-2004
C;Date: 11-Jan-1991 #sequence revision 15-Aug-1996 #text_change 09-Jul-2004
C;Accession: A36106; A30878; A36523; S02173; I54172; I68505
R;Pierce, R.A.; Deak, S.B.; Stolle, C.A.; Boyd, C.D.
Biochemistry 29, 9677-9683, 1990
                                                                                                                                      A; Title: Heterogeneity of rat tropoelastin mRNA revealed A; Reference number: A36106; MUID:91104868; PMID:1702999
                        A; Molecule type: mRNA
A; Residues: 1-864 < PIE>
   A; Cross-references:
                                                                                                   A; Accession: A36106
   UNIPROT: Q99372;
   GB:M60647;
   GB:J05292;
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NID:g207444; PIDN:AAA42269.1;
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A;Cross-references: GB:004035; NID:9207442; PIDN:AAA42268.1; PID:9207443
R;Franzblau, C.; Pratt, C.A.; Faris, B.; Colannino, N.M.; Offner, G.D.; Mogayzel Jr., P.
J. Biol. Chem. 264, 15115-15119, 1989
A;Title: Role of tropoelastin fragmentation in elastogenesis in rat smooth muscle cells.
A;Reference number: A36523; MUID:89359327; PMID:2768256
A;Accession: A36523
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A;Title: Rat tropoelastin is synthesized from a 3.5-kilobase mRNA. A;Reference number: A30878; MUID:88330868; PMID:2971041
A;Accession: A30878
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S59623
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Arch. Biochem. Blophys. 268, 551-558, 1989
Arch. Biochem. Blophys. 268, 551-558, 1989
A;Title: Characterization of rat heart tropoelastin.
A;Reference number: S02173; MUID:89117149; PMID:2913947
A;Accession: S02173
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A;Status: preliminary
A;Molecule type: mRNA
                                                                    R;Mauch, J.C.; Sandberg, L.B.; Roos, P.J.; Jimenez, Matrix Biol. 14, 635-641, 1994
A;Title: Extensive alternate exon usage at the 5' er
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A;Title: Elements of the rat tropoelastin gene asso
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                          A; Reference number: S59623
A; Accession: S59623
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;Residues: 558-864 <RE2>
;Cross-references: GB:M86376; NID:g207459; PIDN:AAA42272.1;
                                                                                                                                                Species: Ovis orientalis aries, Ovis ammon aries, Date: 23-Aug-1996 #sequence_revision 13-Mar-1997, Accession: 859623; A24758
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Residues: 264-533 <RES>
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;Status: preliminary; not compared with conceptual
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Keywords: alternative splicing; extracellular matrix;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cross-references: GB:M86372; NID:g207455;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Status: translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Status: not compared with conceptual translation
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                                                                                                                                                                                                                                                                                                                                                        663
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                                                                                                                                                                                                                                                                                                                                                                                                     EGVRRSLSPELREGDPSSSQHLPSTPSSPRVPGALAAAK 101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  49.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 249.5; DB 1;
Pred. No. 7.8e-13;
3; Mismatches 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene associated with alternative splicing.
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                                                                              end
                                                                                                                                                                        (domestic sheep)
#text_change 09-Jul-2004
translation
                                                                                                                           F.; Christiano, A.M.; Deak,
                                                                            of.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            662
                                                                                                                              S.B.;
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49

GV-GAGVPGFGAGADEGVRRSLSPELREGDPSSSQHLPSTPSSPRVPGA--LAAAK 101

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RESULT 6
A26601
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C;Date: 05-Oct-1988 #sequence revision 26-Jul-1996
C;Accession: A26601; A30795; A27264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R;Yoon, K.; Davidson, J.M.; Boyd, C.; May, M.; LuValle, Arch. Biochem. Biophys. 241, 684-691, 1985
A;Title: Analysis of the 3' region of the sheep elastin A;Reference number: A24758; MUID:85305763; PMID:3839997
                                                                                                                                                                                                                                                                                                                                                                          A;Cross-references: GB:M21880; NID:g212741; PIDN:AAA49082.1; PID:g212742 R;Tokimitsu, I.; Tajima, S.; Nishikawa, T.; Tajima, M.; Fukasawa, T. Arch. Biochem. Biophys. 256, 455-461, 1987 A;Title: Sequence analysis of elastin cDNA from chick aorta and tissue-sp. A;Reference number: A27264; MUID:87297534; PMID:3502711
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F;760-765/Disulfide bonds:
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A; Residues: 655-669,671-716,732-770 < YOO>
밁
                                         á
                                                                                                                                                                                                                                                                                      A;Molecule type: mRNA
A;Residues: 491-569,'G',571-604,'A',606-643,'A',645-687,'R',689-700,'R',702-784 <TOK>
A;Cross-references: GB:M18633; NID:G211742; PIDN:AAA48761.1; PID:G211743
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Baule, V.J.; Foster, J.A.
Biochem. Biophys. Res. Commun. 154, 1054-1060, 1988
A;Title: Multiple chick tropoelastin mRNAs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Residues: 1-212,237-524,535-784 <BR2>
A;Cross-references: UNIPROT: P07916; GB: M15889; NID: g212803;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Reference number: A26601; MUID:87242320;
A;Accession: A26601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A, Title: Repeating structure of chick tropoelastin revealed A, Reference number: A26601; MUID:87242320; PMID:3593675
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Bressan, G.M.; Argos, P.; Stanley, K.K.
Biochemistry 26, 1497-1503, 1987
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N; Alternate names: tropoelastin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               elastin precursor -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Molecule type: mRNA
A; Residues: 1-770 < MAU>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Molecule type: mRNA
A;Residues: 85-784 <BAU>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Reference number: A30795;
A;Accession: A30795
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                            A;Accession: A27264
                                                                                                                                                                              ;Reywords: alternative splicing; extracellular matrix; glycoprotein; 1.24/Domain: signal sequence (fragment) #status predicted <SIG>;25-212,237-524,535-784/Product: elastin #status predicted <MAT>;731-779/Disulfide bonds: #status predicted
                                                                                                                                                                                                                                                                    Superfamily: elast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Superfamily: elastin
                                                                                          Matches
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                                                                                                                 Similarity
AAKFAPGGVGALGGV------GDLGGA-GIPGVGGGVGPAAAAL-GAKAAAK
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                                              AAAAKSAAKVAAKAQLR-----
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                                                                                          Conservative
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                                                                                                               30.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MUID:88309083; PMID:2841924
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                                                                                                               Score 154.5; DB 2
Pred. No. 3.1e-05;
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Pred. No. 2.
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                                                                                               Mismatches
                                -AAAGL---GAGIPGLGV-GVGVPGLGV-GAGVPGL 48
                                                                                                                                                                                                                                                                                                                                                                                   from chick aorta and tissue-specific transcr PMID:3502711
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                                                                                                                                    Length 784;
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Lab. Invest. 51, 605-623, 1984
A;Title: Biology of disease: Elastin: Relation of protein and A;Reference number: I45885; MUID:85059254; PMID:6150137
A;Accession: I45885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          elastin - bovine (fragment)
C;Species: Bos primigenius taurus (cattle)
C;Date: 15-Oct-1996 #sequence_revision 15-Oct-1996 #text_change 09-Jul-2004
C;Accession: I45885
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Theileria annulata
C;Date: 22-Apr-1993 #sequence_revision 18-Nov-1994 #text_change 09-Jul-2004
C;Accession: A45560
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A; Residues: 1-76 < ROS>
N;Alternate names: mastermind protein C;Species: Drosophila melanogaster C;Date: 07-Sep-1990 #sequence_revisio
                                                                       RESULT
A33106
                                                                                                                                                                                                                                                                               Matches
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                                                      neurogenic locus mam protein - fruit fly (Drosophila melanogaster)
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                                                                                                                                                                                                                                                      1 VAAAAKSAAKVAAKAQLRAAAGL------GAGIPGLGV-GVGVPGLGV-GAGVPG
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#sequence_revision 07-Sep-1990 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                         25.9%;
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                                                                                                                                                                                                                                                                                                           Score 130; DB 2;
Pred. No. 0.0033;
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                                                                                                                                                                                                                                                                                                                            Length 907
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A; Experimental source: strain pasc R; Simpson, A.J.G.; Reinach, F.C.; Arruda, P.; Abreu, F.A.; Acencio, M.; Alvarenga, R.; Pariones, M.R.S.; Bueno, M.R.P.; Camargo, A.A.; Camargo, L.E.A.; Carraro, D.M.; Carrer, I as-Neto, E.; Docena, C.; El-Dorry, H.; Facincani, A.P.; Ferreira, A.J.S. submitted to GenBank, June 2000

A; Authors: Ferreira, V.C.A.; Ferro, J.A.; Fraga, J.S.; Franca, S.C.; Franco, M.C.; Frohw J.D.; Junqueira, M.L.; Kemper, E.L.; Kitajina, J.P.; Krieger, J.E.; Kuramae, B.E.; Lalguchado, M.A.; Madeira, M.M. Madeira, H.M.F.; Marrino, C.L.; Marques, M.V.; Martins, E.M. F.; Matsukuma, A.Y.; Menck, C.F.M.; Miracca, E.C.; Miyaki, C.Y., F.G.; Nunes, L.R.; Oliveira, M.A.; de Oliveira, M.C.; de Oliveira, R.C.; Palmieri, D.? Rofigues, V.; Rosa, A.J. de M.; de Rosa Jr. V.E.; de Sa, R.G.; Santelli, R.V.; Sawasak, A; Authors: da Silva, A.C.R.; da Silva, F.R.; da Silva, A.M.; Silva Jr., W.A.; da Silvai, M.; Tsuhako, M.H.; Vallada, H.; Van Sluys, M.A.; Verjovski-Almeida, S.; Vettore, A.L.; A.R.; Salvasia, R.C.; R.S.; Santelli, R.V.; Sawasak, R.S.; Vettore, A.S.; Van Sluys, M.A.; Verjovski-Almeida, S.; Vettore, A.L.; A.Reference number: A59328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R;Smoller, D.; Friedel, C.; Schmid, A.; Bettler, D.; Lam, L.; Genes Dev. 4, 1688-1700, 1990
A;Title: The Drosophila neurogenic locus mastermind encodes a A;Reference number: A36391; MUID:91065516; PMID:1701150
A;Accession: A36391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Title: The genome sequence of the plant pathogen Xylella fastidiosa. A;Reference number: A82515; MUID:20365717; PMID:10910347 A;Note: for a complete list of authors see reference number A59328 below A;Accession: E82734 A;Status: preliminary A;Molecule type: DNA A;Residues: 1-127 <SIM> A;Resi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hypothetical protein XF1025 [imported] - Xylella fastidiosa (strain 9a5c) C;Species: Xylella fastidiosa C;Date: 18-Aug-2000 #sequence_revision 20-Aug-2000 #text_change 09-Jul-2004 C;Accession: E82734
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A;Cross-references: UNIPROT:P21519; GB:X54251;
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22 GLGAGIPGLGVGVGVPGLGVGAGVPGLGVGAGVPGFGAGADEGVRRSLSPELREGDPSS
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33.9%; Pred. No. 0.061;
tive 13; Mismatches
                                                                                                                                                                                        Score 112.5; DB Pred. No. 0.015; 7; Mismatches
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C;Accession: T29074

R;Redenbach, M.; Kieser, H.M.; Denapaite, D.; Eichner, A.; Cullum, J.; Kinashi, Mol. Microbiol. 21, 77-96, 1996

A;Title: A set of ordered cosmids and a detailed genetic and physical map for th A;Reference number: Z20556; MUID:97000351; PMID:8843436

A;Accession: T29074
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A; Residues: 1-532 < SEE>
A; Cross-references: UNIPROT:069975;
A; Experimental source: strain A3(2)
C; Genetics:
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C;Species: Streptomyces coelicolor
C;Date: 05-Nov-1999 #sequence_revision 05-Nov-1999 #text_change 09-Jul-2004
C;Accession: T35119
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Plant Mol. Biol. 35, 969-972, 1997
A;Title: A novel gene of tomato preferentially expressed
A;Reference number: Z15238; MUID:98088022; PMID:9426616
A;Accession: T04143
A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                          hypothetical protein SCIC2.25c - Streptomyces coelicolor C;Species: Streptomyces coelicolor C;Date: 22-Oct-1999 #sequence_revision 22-Oct-1999 #text
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A; Residues: 1-505 < KI
                    A; Molecule type: DNA
A; Residues: 1-1329 <
                                                     A; Status: preliminary; translated
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Best Local S
Matches 35
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Matches 32
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Cross-references: UNIPROT:086525; EMBL:AL031124; NID:e1312893; PID:e1312918; PIDN:CAA1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1-505 <KIY>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAKSAAKVAAKAQ-----LRAAAGIGAGIPGIGVGVGVPGIGVGAGVPGIGVGA 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FGAGA-DEGVRRSLSPELREGDPSSSQHLPSTP 88
                                                                                                                                                                                                                                                                                                                                        GTGPGTGPGA--GGRRSAPRRQAAREPAARTYARALPVVP 133
                                                                                                                                                                                                                                                                                                                                                                                 GV-PGFGAGADEGVRRSLSPELREGDPSSSQHLPSTPSSP 91
                                                                                                                                                                                                                                                                                                                                                                                                                         ASEPAAGTSTKAHTARVGAGQGRGLHEGAGQGSGMAGSGSGTG-PGSGTG---PGTGTGP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SGFGAVGSGLSKAGKFMGRTFTGSSKKNGSSTP 489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                      <RED>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Parkhill, J.;
Library, March
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 109; DB Pred. No. 0.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 109.5; DB Pred. No. 0.088; 7; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         from
                                                          from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL: AL022268; PIDN: CAA18336.1; GSPDB: GN00070; SCOEI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                          GB/EMBL/DDBC
                                                                                                                                                                                            22-Oct-1999 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-Apr-1999 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Barrell, B.G.; Rajandream, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   37;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         fruit
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18;
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C;Accession: G86252

R;Theologis, A.; Ecker, J.R.; Palm, C.J.; Federspiel, N.A.; Kaul, S.; White, O.; Alonso Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; Dewar, K. Chin, C.W.; Chung, M.K.; Conn, L.; Conway, A.B.; Conway, A.R.; Creasy, T.H.; Dewar, K. ansen, N.F.; Hughes, B.; Huizar, L.

Nature 408, 816-820, 2000

A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khaykin, B.; Kim, C. A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Maiti, R.; Marziali Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.; Z.A.; Luros, J.S.; Maiti, R.; Marziali Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.; Southwick, A.M.; Sun, H.; Tallon, ker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.

A;Title: Sequence and analysis of chromosome 1 of the plant Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             glycine-rich protein - common sainfoin
C;Species: Onobrychis viciifolia (common sainfoin)
C;Date: 21-May-1999 "sequence_revision 21-May-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       T08002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hypothetical protein [imported] - Arabidopsis thaliana
C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 09-Jul-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ś
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            submitted to the EMBL Data Library, October 1997
A,Description: A cDNA coding for a glycine-rich protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
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                                                                     A; Map position:
C; Superfamily:
                                                                                                                                   A; Cross-references:
                                                                                                                                                      A;Residues:
                                                                                                                                                                          A; Molecule type: DNA
                                                                                                                                                                                                                      A;Reference number: A86141; MUID:21016719; PMID:11130712
A;Accession: G86252
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: mRNA
A; Residues: 1-151 < JOS>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Status: preliminary; translated from GB/EMBL/DDBC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Accession: T08002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Reference number: 216281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R; Joseph,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Note:
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                                                                                                                 Genetics:
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Query Match
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                                                                                                                                                      1-108 <STO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 VAAAAKSAAKVAAKAQLR--------AAAGLGA-GIP--GLGVGVGVGVPGLGVGAGV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 AAAKSAAKVAAKAQLRAAAGLGAGIPGLGVGVGVFGLGVGAGVPG---LGVGAG-----VPG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PGL-GVGAGVPGFGAGADEG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAAQAAAGVAAAR---RGGQGGGALAGGGAFGGGSGGGALGAGAPGGRLGAGQGPRAALPG
                                                                glycine-rich RNA-binding protein; ribonucleoprotein repeat homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                   UNIPROT:065397; GB:AE005172; NID:g3157948; PIDN:AAC17631.1; GSPDB:G
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  20.7%;
49.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10;
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Pred. No. 0.082;
  Score 103.5; DB Pred. No. 0.067;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 105; DB:
Pred. No. 0.48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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                         2:
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Title:
Perfect score:
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Maximum DB
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    Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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Copyright (c) 1993.25 2005 Compugen Ltd
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oa); NAS. conferring	itel Homo by J.,	08pzf2 Q25946 Q6z2m1 Q9cz2m1 Q9c5790 P215190 Q9pek3 Q9Bjw2 Q9bjw2 Q9bjw2 Q9bjw2 Q9bjw2 Q9bjw2 Q9bjw2 Q9bjw2 Q9bjw2 Q9bjw2
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WAIDLINE-22737999; PubMed=12853948; DOI=10.1038/nature01782;
WAIR Hillier L.W., Fulton R.S., Fulton L.A., Graves T.A., Pepin K.H.,
WAWA Wagmer-McPherson C., Layman D., Maas J., Jaeger S., Walker R.,
WAIR K., Sekhon M., Becker M.C., O'Laughlin M.D., Schaller M.E.,
WAIR K., Sekhon M., Becker M.C., O'Laughlin M.D., Schaller M.E.,
WAIR M.D., Delehaunty K.D., Miner T.L., Nash W.E., Cordes M., Du H.,
WAIR M.D., Delehaunty K.D., Miner T.L., Nash W.E., Cordes M., Du H.,
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WAIR M.D., Delehaunty K.D., Miner T.L., Nash W.E., Cordes M., Du H.,
WAIR M.D., Belcki L., Scott K., Holmes A., Harkins R., Harris A.,
WAIR M.D., Harkins R., Harris A.,
WECTONG C.M., Hou S., Tomlinson C., Dauphin-Kohlberg S.,
WECTONG C.M., Hou S., Tomlinson C., Dauphin-Kohlberg S.,
WECTONG C.M., Hou S., Tomlinson C., Dauphin-Kohlberg S.,
WECTONG C.M., Hou S., Tomlinson D., Maupin R., Stromatt C.,
WAIR M., Harris A., Walling T., Woessner J.P.,
Wendl M.C., Yang S.P., Schultz B.R., Wallis J.W., Spieth J.,
Wallie P., Miller N., Johnson D., Murray J., Woessner J.P.,
Wendl M.C., Yang S.P., Schultz B.R., Wallis J.W., Spieth J.,
Wallifton S.W., Chissoe S.L., Marra M.A., Raymond C., Haugen E.,
WAIR M., Waller W., Zhou Y., James R., Phelps K., Iadanoto S., Bubb K.,
WAIR M., Waller W., Walliam D., Wallanoto S., Bubb K.,
WAIR M., Waller W., Walliam S., Wallanoto S., Bubb K.,
WAIR M., Waller W., Walliam S., Flicek P., Bork P., Suyama M.,
WAIR M., Waller W., Waller B., Flicek P., Bork P., Suyama M.,
Waller W., Waller M., Waller B., Flicek P., Bork P., Suyama M.,
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Q75MU5;
Q5-JUL-2004
05-JUL-2004
05-JUL-2004
                                                                                                                                                                     EMBL; AC005056; AAS07435.1; -
GO; GO:0005578; C:extracellular matr
GO; GO:0005201; F:extracellular matr
InterPro; IPR001451; Hexapep transf.
InterPro; IPR003979; tropoelastin.
PRINTS; PR01500; TROPOELASTIN.
                                                                                         Hypothetical SEQUENCE 75
                                                                                                                                                                                                                                                                                                                                                       Submitted (JAN-2004) to the EMBL/GenBank/DDBJ
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nature 424:157-164(2003).
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   Score
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                                                                                         2B24F955D8360738 CRC64;
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   501;
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   L; DB 2;
. 8.6e-28;
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                           Length
                                                                                                                                                                                                                                                                    constituent; IEA.
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; Homo.
                                                                                                                                                                                                                                                                                                   IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Suyama M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isak A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               <u>.</u>
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RESULT 4
Q95WFS
ID Q95W
AC Q95W
AC Q95W
DT 01-M
DT 01-M
DT 01-M
DT 01-M
CO Elas
GN Name
OS Homo
OC Euka
OC WClm
RN Mamm
OC WClm
RN SEQU
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Q9UMF5;
01-MAY-2000
01-MAY-2000
01-MAR-2004
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EMBL; U93035; AAB65621.1; JOINED.
EMBL; U93035; AAB65621.1; JOINED.
EMBL; U93036; AAB65621.1; JOINED.
GO; GO:0005578; C:extracellular matrix (sensu Metazoa); IEA.
GO; GO:0005201; F:extracellular matrix structural constituent; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Li D.Y., Toland A.E., Boak B.B., 1
Morris C.A., Keating M.T.;
"Elastin point mutations cause an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
Mammalia; Eutheria;
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01-JAN-1998 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
                                    Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                   Elastin (Fragment).
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              NCBI_TaxID=9606;
[1]
                                                             Homo sapiens (Human)
                                                                           Name=ELN;
                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR003:
PRINTS; PR01500;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              supravalvular aortic stenosis.'
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 SEQUENCE FROM N.A
                                                                                                                                                                                                                                                                                                                                                                                                       PROSITE; PS00101; HEXAPEP TRANSFERASES; UNKNOWN_1
                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR001451; Hexapep_transf.
InterPro; IPR003979; tropoelastin.
PRINTS; PR01500; TROPOELASTIN.
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                                                                                                                                                                                                                                                                                                                                                                   635 AA;
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Primates;
                                    Chordata;
Primates;
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Last annotation updat
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                                       Catarrhini;
                                                 Craniata;
                                                                                                                                                      PRT;
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Atkinson D.L., Ensing G.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           obstructive vascular disease,
                                                                                                                                                                                                                                                                                                                                 No.
                                                                                                                                                                                                                                                                                                                                            493;
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                                                   Vertebrata; Euteleostomi;
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                                       Hominidae;
                                                                                                  update)
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Best Local S
Matches 68
                                                                                                         Query Match
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Matches 68
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                                                                                                                                                                                                                                                                                                                                                                                                              01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Hypothetical protein PSEC0254.
                                                                                                                                                              GO; GO:0005578; C:extracellular matrix (sensu Metazoa); IEA.
GO; GO:0005201; F:extracellular matrix structural constituent;
InterPro; IPR001451; Hexapep_transf.
InterPro; IPR003979; tropoelastin.
PRINTS; PRO1500; TROPOELASTIN.
PROSTITS; PRO1500; TROPOELASTIN.
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EMBL; U63721; AAC13884.1; -.

InterPro; IPR001451; Hexapep_transf.

PROSITE; PS00101; HEXAPEP_TRANSFERASES; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Identification of genes from a 500-kb commonly deleted in Williams syndrome p Genomics 36:328-336(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=96411691; PubMed=8812460; DOI=10.1006/geno.1996.0469;
Osborne L.R., Martindale D.W., Scherer S.W., Shi X.-M., Huizenga J.,
Heng H.H.Q., Costa T., Pober B., Lew L., Brinkman J., Rommens J.,
Koop B.F., Tsui L.-C.;
                                                                                                                                                                                                                                                                                                                                                                                      Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q8NBI4;
01-OCT-2002
                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Q8NBI4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Comparative genomic sequence analysis of the Williams syndrome region (LIMK1_RFC2) of human chromosome 7q11.23.";
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                         ADEGVRRSLSPELREGDPSSSQHLPSTPSSPRVPGALAAAK 101
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Pred. No. 2.
                                                                                                          Score 304.5; DB 2
Pred. No. 6.3e-14;
D; Mismatches 0
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Nakamura Y.
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Homo sapiens (Human).
Bukaryota; Metazoa; C
Mammalia; Eutheria; F
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Q14235;
Q1-NOV-1996
Q1-DEC-2001
Q1-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                 GO; GO:0005578; C:extracellular matrix (sensu Metazoa); GO; GO:0030023; F:extracellular matrix constituent confe InterPro; IPR001451; Hexapep_transf. InterPro; IPR001379; tropoelastin.
PRINTS; PR01500; TROPOELASTIN.
PROSITE; PS00101; HEXAPEP_TRANSPERASES; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rosenbloom J., Ornstein-Goldstein N., "Structure of the 3' region of the human el of Alu repetitive sequences and few coding Connect. Tissue Res. 16:197-211(1987).
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687 AA; 59529 MW; 864068C4C8E9F88F CRC64;
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01-CCT-2003 (TrEMBLrel. 2:
01-MAR-2004 (TrEMBLrel. 2:
                                                                                                                                                                                                                                                                                                                                                            Poustka A., Albert R., Moosmayer P., Schupp I., Wellenre Mewes H.W., Weil B., Amid C., Osanger A., Fobo G., Han Mewes H.W., Weil B., Amid C., Osanger A., Fobo G., Han Mewes H.W., Weil B., Chools to the EMBL/GenBank/DDBJ databases.
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Hypothetical protein DKFZp686021208 (Fragment).
                                                                                                                                                                                                                                                                                                          PROSITE; PS00101; HEXAPEP_TRANSFERASES; UNKNOWN_1.
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Lauber J., Bahr A.,
Han M., Wiemann S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota;
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PROSITE; PS00101; HEXAPEP_TRANSFERASES;
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TISSUE=Human rectu
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Primates;
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Pred. No. 6.8e-14;
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No. 6.6
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elastin gene (ELN).";
J. Biol. Chem. 274:98
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Rosenbloom J.C., Peltonen L., Rosenbloom J.;
                                                                                                                                                            Urban Z., Michels V.V., Thibodeau S.N., Davis E.C., Bonnefont J.-P., Munnich A., Eyskens B., Gewillig M., Devriendt K., Boyd C.D.; "Isolated supravalvular aortic stenosis: functional haploinsufficiency of the elastin gene as a result of nonsense-mediated decay."; Hum. Genet. 106:577-588(2000).

-!- FUNCTION: Major structural protein of tissues such as aorta and nuchal ligament, which must expand rapidly and recover completely.
-!- SUBUNIT: The polymeric elastin chains are cross-linked together into an extensible 3D network.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=89009960; PubMed=3171221; Fazio M.J., Olsen D.R., Kauh B.A., Baldwin C.T., Indik Z., Cornstein-Goldstein N., Yeh H., Rosenbloom J., Uitto J.; "Cloning of full-length elastin cDNAs from a human skin fibroblast recombinant cDNA library: further elucidation of alternative splic utilizing exon-specific oligonucleotides."; J. Invest. Dermatol. 91:458-464(1988).
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P15502; Q14233; Q14238;
01-APR-1990 (Rel. 14, Created)
01-APR-1990 (Rel. 14, Last sequence up
25-OCT-2004 (Rel. 45, Last annotation
Elastin precursor (Tropoelastin).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE-Hippocampus, and Placenta;
MEDLINE-96291399; PubMed-8689688; DOI=10.1016/S0092-8674(00)80077-X;
MEDLINE-96291399; DOI=10.1016/S0092-8674(00)8007-X;
MEDLINE-96291399; DOI=10.1016/S0092-8674(00)8007-X;
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MEDLINE-9
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"Isolation and characterization of human elastin
associated variation in elastin gene expression i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE OF 603-730 FROM N.A. TISSUE-Hippocampus, and Placenta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fazio M.J.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PubMed=10942104;
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                                                            SUBCELLULAR LOCATION: Extracellular ALTERNATIVE PRODUCTS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Invest. 58:270-277(1988).
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J., Uitto J.;
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EMBL; M17282; AAC98394.1; J

EMBL; M17265; AAC98394.1; J

EMBL; M17266; AAC98394.1; J

EMBL; M17266; AAC98394.1; J

EMBL; M17267; AAC98394.1; J

EMBL; M17270; AAC98394.1; J

EMBL; M17271; AAC98394.1; J

EMBL; M17273; AAC98394.1; J

EMBL; M17273; AAC98394.1; J

EMBL; M17275; AAC98394.1; J

EMBL; M17276; AAC98394.1; J

EMBL; M17277; AAC98394.1; J

EMBL; M17278; AAC98394.1; J

EMBL; M17280; AAC98394.1; J

EMBL; M17280; AAC98394.1; J

EMBL; M36860; AAA53190.1; -

EMBL; M36860; AAA53190.1; -

EMBL; M35603; CAAA53190.1; -

EMBL; M35603; CAAA53190.1; -
              PRINTS; PR01500; TROPOELASTIN.
Alternative splicing; Repeat; Signa Williams-Beuren syndrome.
SIGNAL 1 26
                                                                                                                                                           8000000 MIM
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                                                                                                                                                                                                                                                                                                                                                                                                      Genew; HGNC:3327; ELN.
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HSSP; P50099; 1ZFC
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                                                                                                                                                      GO:0005578; C:extracellular matrix; TAS.
GO:0005615; C:extracellular space; TAS.
GO:0005201; F:extracellular matrix structural
GO:0008283; P:cell proliferation; TAS.
GO:0008015; P:circulation; TAS.
GO:0009807; F:organogenesis; TAS.
GO:0009807; P:respiratory gaseous exchange; TF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     laxa [MIM:123700]. Cutis laxa is a rare connective tissue disorde characterized by loose, hyperextensible skin with decreased resilience and elasticity leading to a premature aged appearance. The skin changes are often accompanied by extracutaneous manifestations, including pulmonary emphysema, bladder diverticula, pulmonary artery stenosis and pyloric stenosis. DISEASE: Haploinsufficiency of DISEASE: Haploinsufficiency of DISEASE: Haploinsufficiency of cardiovascular and musculo-skeletal abnormalities observed in williams-Beuren syndrome (WBS) [MIM:194050]. WBS is a rare developmental disorder and a contiguous gene deletion syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SWISS-PROT entry is copyright. It is produced need the SWiss Institute of Bioinformatics and the European Bioinformatics Institute. There are no rest buropean Bioinformatics Institutions as long as its content by non-profit institutions as long a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   developmental disorder and a contiguous gene deletion syndrome involving genes from chromosome band 7q11.23.

DISCASE: Defects in ELN are these of supravalvular aortic stenosis (SVAS) [MIM:185500]. SVAS is a congenital narrowing of the ascending aorta which can occur sporadically, as an autosomal dominant condition, or as one component of Williams-Beuren
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Q6ZWJ6;
05-JUL-2004
05-JUL-2004
05-JUL-2004
                                01-OCT-2002
01-OCT-2002
01-MAR-2004
                                                                  Q8N2G0;
                                                                                                                                                                                                                                                                                                                                                                                                   Tanigami A., Fujiwara T., Shibahara T., Goto Y., Hirao M., Shimizu Wakebe H., Ono T., Hishigaki H., Watanabe T., Ozaki K., Sugiyama T. Irie R., Otsuki T., Sato H., Wakamatsu A., Ishii S., Yamamoto J., Isono Y., Kawai-Hio Y., Saito K., Nishikawa T., Kimura K., Yamashita H., Matsuo K., Nakamura Y., Sekine M., Kikuchi H., Kanda Wagatsuma M., Murakawa K., Kanehori K., Takahashi-Fujii A., Oshima Sugiyama A., Kawakami B., Suzuki Y., Sugano S., Nagahari K.,
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05-JUL-2004 (TrEMBLrel. 27, Last sequence update)
05-JUL-2004 (TrEMBLrel. 27, Last annotation update)
Hypothetical protein FLJ16246.
Homo sapiens (Human).
                                                                                                                                                                                                                                                                                 Masuho Y., Nagai K., Isogai T.;
Submitted (JUL-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; AK122731; BAC85506.1; -.
GO; GO:0005578; C:extracellular matrix (sensu Metazoa);
GO; GO:0005201; F:extracellular matrix structural consti
InterPro; IPR001451; Hexapep transf.
InterPro; IPR003979; tropped_astin.
PRINTS; PRO1500; TROPOELASTIN.
PRONSITE; PS00101; HEXAPEP_TRANSFERASES; UNKNOWN 1.
PROSITE; PS00101; HEXAPEP_TRANSFERASES; UNKNOWN 1.
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         Hypothetical protein PSEC0191.
Homo sapiens (Human)
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Chordata; Craniata;
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1; Mismatches 0
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Missing (in isoform 2).
/FTId=VSP_004243.
, AB06D15BA567AE46 CRC64;
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ugiyama T.,
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RA Nagahari K., Sugano S., Isogai T.;

Submitted (MAR-2002) to the EMBL/GenBank/DDBJ databases.

REMBL; AK075494; BAC11651.1;

RGO; GO:0005578; C:extracellular matrix (sensu Metazoa); IEA

RGO; GO:0005201; F:extracellular matrix structural constitue.

RINTERPRO; IPR001451; Hexapep transf.

RINTERPO; IPR003979; tropoelastin.

RRINTS; PR01500; TROPOELASTIN.
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Best Local S
Matches 67
                                                                                                 Query Match
Best Local Similarity
Matches 67; Conser
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015337;
01-JAN-1998 (TrEMBLrel. (
01-JAN-1998 (TrEMBLrel. (
01-MAR-2004 (TrEMBLrel. (
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SEQUENCE
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EMBL; U93034; AAB65620.1; JOINED.
EMBL; U93035; AAB65620.1; JOINED.
EMBL; U93036; AAB65620.1; JOINED.
GO; GO:0005578; C:extracellular matrix (sensu Metazoa); IEA.
GO; GO:0005201; F:extracellular matrix structural constituent;
InterPro; IPR001451; Hexappe_transf.
InterPro; IPR003979; tropoelastin.
PRINTS; PR01500; TROPOELASTIN.
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MEDLINE=97358574; PubMed=9215670; DOI=10.1093/hmg/6.7.1021;

MEDLY., Toland A.E., Boak B.B., Atkinson D.L., Ensing G.J.
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Mammalia; Eutheria;
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Hum. Mol. Genet. 6:1021-1028(1997).
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l Similarity 66.3%;
67; Conservative
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                                VAAAAKSAAKVAAKAQLKAAAGLGAGIPGIGVGVGVGVGVGAGVPGLGVGAGVPGFGAG
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Pred. No. 2.2e-13;
0; Mismatches 1
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Nakamura Y.,
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COLL Res. 7:235-247(1
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01-NOV-1996
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation updat
Elastin-cBEL1; NCBI gi: 163003 (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           splicing.";
Coll. Relat
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"Sequence variation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NON_TER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ornstein-Goldstein N., Indik Z., Sheppard oom J.C., Cicila G., Yoon K., Rosenbloom J ce variation of bovine elastin mRNA due to
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                    VAAAAKSAAKVAAKAQLRAAAGLGAGIPGLGVGVGVGVDGLGVGAGVPGLGVGAGVPGFGAG
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Pred. No. 8.5e-12;
1; Mismatches 6
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structural constituent;
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Best Local S
Matches 61
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"Sequence variation of bovine elast
splicing.";
Coll. Relat. Res. 7:235-247(1987).
EMBL; M19372; AAA30501.1; JOINED.
EMBL; M19366; AAA30501.1; JOINED.
EMBL; M19367; AAA30501.1; JOINED.
EMBL; M19367; AAA30501.1; JOINED.
EMBL; M19369; AAA30501.1; JOINED.
EMBL; M19369; AAA30501.1; JOINED.
EMBL; M19370; AAA30501.1; JOINED.
EMBL; M19371; AAA30501.1; JOINED.
EMBL; M22772; AAA30501.1; JOINED.
EMBL; M22773; AAA30501.1; JOINED.
EMBL; M22773; AAA30501.1; JOINED.
EMBL; M22773; AAA30501.1; JOINED.
EMBL; M22773; AAA30501.1; JOINED.
EMBL; M22774; AAA30501.1; JOINED.
EMBL; M22775; AAA30501.1; JOINED.
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Q28098;
Q1-NOV-1996 (TrEMBLrel. 01, Created)
Q1-NOV-1996 (TrEMBLrel. 01, Last sequence update)
Q1-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Elastin; NCBI gi: 163002 (Fragment).
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GO; GO:0005201; F:extracellular matrix structural constituent;
INTERPRO; IRRO3979; troppelastin.
PRINTS; PR01500; TROPOELASTIN.
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SEQÜENCE 666 AA; 56435 MW; BCB5E62632BE1B71 CRC64:
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MEDLINE=85280426; PubMed=2992576;

Cicila G., May M., Ornstein-Goldstein N.,

Yeh H.S., Rosenbloom J., Boyd C., Rosenblo
"Structure of the 3' portion of the boving
Biochemistry 24:3075-3080(1985).
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01-JUN-2003 (TrEMBLrel.
Blastin-cBEL3; NCBI gi:
                                                                                                                                      Bos taurus (Bovine).
Eukaryota; Metazoa;
Mammalia; Eutheria;
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Bovinae; Bos.
  SEQUENCE FROM N.A. MEDLINE=85280426; PubMed=2992576;
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                                                                                                                                                                    Chordata; Craniata; Vertebrata;
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24, Last annotation update)
163005 (Fragment).
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Pred. No. 8.6e-12;
1; Mismatches 6
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Rosenbloom J
                                                                                                                                            Ruminantia;
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minantia; Pecora; Bovidae;
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R EMBL; M19370; AAA30498.1; JOINED.
R EMBL; M19370; AAA30498.1; JOINED.
R EMBL; M27771; AAA30498.1; JOINED.
R EMBL; M22772; AAA30498.1; JOINED.
R EMBL; M22773; AAA30498.1; JOINED.
R EMBL; M22773; AAA30498.1; JOINED.
R EMBL; M22773; AAA30498.1; JOINED.
R EMBL; M22774; AAA30498.1; JOINED.
R EMBL; M22775; AAA30498.1; JOINED.
R EMBL; M22775; AAA30498.1; JOINED.
R EMBL; M22775; AAA30498.1; JOINED.
R EMBL; M22988; AAA30498.1; JOINED.
R GO; GO:0005201; P:extracellular matrix (sensu Metazoa); IEA GO; GO:0005201; F:extracellular matrix structural constitue interpro; IPR003979; troppoelastin.
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Coll. Relat. Res. 7:235-247
EMBL; M19372; AAA30498.1; JUNE 1, AAA
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SEQUENCE
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Rosenbloom J.C., Cicila G., Yoon K.,
                                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPR003979; tropoelas
PRINTS; PR01500; TROPOELASTIN.
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543
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                                                                                                                                        1 VAAAAKSAAKVAAKAQLRAAAGLGAGIPGLGVGVGVPGLGVGAGVPGLGVGAGVPGFGAG
                                                                                                                                                                                                                                                         Similarity
                                                                                                             VPAAAKSAAKAAAKAQFRAAAGLPAGVEGUGVGVGVGUFGLGVGVGVEGLGVGAGVPGFGA-
                                                     ADEGVRRSLSPELREGDPSSSOHLPSTPSSPRVPGALAAAK
                                                                                                                                                                                                                                                                                                                                            707
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                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                            AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PubMed=3665402;
Goldstein N., Indik
                                                                                                                                                                                                                                                                                                                                            60346 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of bovine elastin
                                                                                                                                                                                                                                                      54.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               JOINED
                                                                                                                                                                                                                           Score 274.5; | Pred. No. 9.1e
                                                                                                                                                                                                                                                                                                                                         FDFD559BAB34CE33 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Z., Sheppard P., Rosenbloom J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mRNA due to
                                                                                                                                                                                                                                                      .1e-12;
                                                                                                                                                                                                                                                                                          DB
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                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                             constituent;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                alternative
                                                                                                                                                                                                                                 Indels
                                                        101
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                                                                                                                                                                                                                                                                                    707;
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